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## ALIGNMENTS

BASE COUNT ORIGIN		source	FEATURES		JOURNAL	TITLE	AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	AX105267	RESULT 1
expression" 492 a 236 c 230 y 492 t	/organisme"synthetic construct" /db_xref-"taxon:32630" as a Promoter for regulating /note-"Sequence to be used as a Promoter for regulating	11350	Location/Qualifiers	Medicago Inc. (CA)	Patent: WO 0125455-A 1 12-APR-2001;	Promoter for regulating expression of foreign genes	Vezina, L.P. and D'Aoust, M.A.	1 (bases 1 to 1350)	artificial sequence.	synthetic construct	synthetic construct.		AX105267.1 G1:13921417	AX105267	Sequence   from Patent W00125455.	AX105267 1350 bp FNA linear PAT 30-APF-2001		

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Patent: WO 0125455-A 3 12-APR-2001;
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Poichert, T., Tenzelewski, V. and Haehnel, W.
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1 (bases 1 to 326)
Rance, I., Theisen, M. and Gruber, V.
Patent: FP 2791858-A 2 29-SEP-2000;
MERISTEM THERAPEUTICS (FR)
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                                                                                                        Freiburg, Lehrstuhl fuer Biochemic Freiburg, Germany, D-79104
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/db_xref="taxon·364]"
/tissue_type="leaf"
                                      /strain-"Var. Italica"
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Yamada, K., Liu, S.X., Sakano, H., Phame, P.K., Banh, J., Thord, M.K., Goldsmith, A.D., Lee, J.M., Toriumi, M., Yu, B., Hrooks, S., Chao, O., Chen, H., Earlin Neumann, C., Kim, C., Lam, P., Miranda, M., Nargeb, M. Palm, C.J., Shinu, P., Southwick, A., Davis, P.W., Feker, J.R. and Theologis, A.
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Yamada, K., Lita, S.A., Sakano, n., Pham. P.K., Banh. J., Froma, M.K., Yamada, K., Lita, S.A., Sakano, n., Pham. P.K., Banh. J., Froma, M.K., Chao, G., Chao, B., M., Lan, B., Mitanha, M., Nhayen, M., Chen, H., Karlin, Neumann, G., Kim, C., Lam, B., Mitanha, M., Nhayen, M., Palm, C.L., Shirm, P., Southwick, A., Lavis, B.K., Teker, L.K., and Palm, C.L., Shirm, P., Southwick, A., Lavis, B.K., Teker, L.K., and
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Arabidopsis thaliana (strain Columbia) (library: Lambda-YES-R)
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                                            /protein_id="AAA32787.1"
/db_xre1="G1:16b996"
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/translation="MASTSTTVPIESFTGLKASTTKSSATVETQTAAVASPKLTVKS
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YPHNVVEDEDEIPSGVDVAKISMDEQDLLNGAGETYEVALTEPGTYSFYCAPHQGAGM
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Submitted (29 NOV 20 ) Salk Institute Copemic Analysis Laboratory (SIGnAL), Plant Buckery Laboratory, Eme Salk Institute for Biological Studies, Ca910 N. Forrey Pines Road, La Jolla, CA 92037
                                                                                                                                                                                                                                                                                         Southwick, A., Toriumi, M., Yamada, K., Yu, G., Shinozaki, K., Davis, R.W., Theologic, A. and Ecker, J.R.
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Arabidopsis Full Length CDNA'): Scki.m., Narusaka, M.,
Satou, M., Kamiya, A., Sakurai, L., Carbinot, F., Kawai, ...
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Copyright (c) 1993 - 2000 Compugen Ltd
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| STIRST/grightara/geneserg/genesergn-cmbt/NAT984.LAT1*
| STIRST/grightara/geneserg/genesergn-cmbt/NAT986.LAT1*
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ionserdens income suppressor	AAS46311	22	7913	4.7	63	3. 14	
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Tumbur suppressor	AAS46429	es Es	6106		63.8	Ę.	ဂ
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Nucleotide sequenc	AAA96460	7	207	ස ි	116	10	

## ALIGNMENTS

AASO2126 ID AASO

AAS02126;

AAS02126 standard; DNA; 1350 BP

RESULT

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Novel promoter for regulating expression of foreign genes organisms, more specifically in a loaf specific manner in WP1, 2001 266316/27. (MEDI-) MEDICAGO INC 04-OCT-1999; Vezina h, 02-0CT-2000; 2000WO-CAU1144 12-APR-2001. W0200125455-A2 Synthetic Promoter, transgenie plant; monocotyledon; dicotyledon; gymnooperm; ds Promoter #1 for regulating expression of foreign genes 18- JUL-2001 D'AJust M, (first entry) 99us-0157129 in transgenic

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The sequence represents the coding sequence of promoter #2 for regulating expression of foreign genes in transgenic organisms. The promoter is

Claim 1; Page 9; 9pp; English.

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Query Match
Best Local S
Matches 971
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                                                                                                                                                                                                                                                                                                                 useful for regulating the expression of foreign genes in transgenic organisms, particularly plants, e.g. monocotyledons, dicotyledons or gymnosperms, by preparing a transgenic organism using ab expression construct comprising the promoter and an open reading frame (ORF) of
                                                                                                                                                                                                                                                                                                         gene
                                                                                                                                                                                                                                                                                                                                                                      The sequence represents the coding sequence of promoter #3 for regulating expression of foreign genes in transgenic organisms. The promoter is
                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 9; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      organisms, more specifically in a leaf specific manner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel promoter for regulating expression of forelyn genes in transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WP1; 2001-266316/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MEDI-) MEDICAGO INC
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                                                                                                                                                                                                                                                                                Sequence 731 EP;
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Pred. No.
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AAA96458
    Claim 2; Page 56
                                                      Chimeric expression promoter for producing dicotyledomous and monocotyledomous transactic plants comprises a nacteic acid sequence
                                    derived from a promoter of the pea plastocyanin gene
                                                                                                                  WPT: 2000 587662/55
                                                                                                                                                  Rance I, Gruber V.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the present sequence represents the chimeric promoter Mp1092. (he promoter is derived from the petE promoter from pea plastocyanin gene-
by a deletion in 5' of the repeated invert sequences as well as the
as-1 like box borne on a Sphl fragment of 212 bp. The petE promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expression promoter comprising a promoter of the pea plastocyanin dene, or comprising a 3 box operably or functionally linked upstream of a CAAT box. TATA box and transcription initiation site. The chimeric promoters
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Range I, Gruber V,
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                                                                                                                                                                                                                                                                                                                                                                                                                                       W0200056906 A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08 FEB-2001 (first entry)
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                                                                                 (MERL ) MEKISTEM THEKAPEULING
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                                                                                                                                                                                                                                                                                                               08-FEB-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 65; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric expression promoter for producing dicotyledomous and monocotyledomous transgenic plants comprises a nucleic acid sequence
              (MERI-) MERISTEM THERAPEUTICS
                                               22-MAR-1999;
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                                                                                                              28-SEP-2000
                                                                                                                                              W0200056906-A1
                                                                                                                                                                                 Pisum sativum
                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                plastocyanin
                                                                                                                                                                                                                                                petE promoter; chimeric promoter;
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                                                                                                                                                                                                                                gene
                                               99FR-0003635
                                                                                                                                                                                                                                promoter; ss
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82 8%;
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                                                                                                                                                                                                                                                  MPr1096;
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                                                                                                                                                                                                                                                                                                                                       promoter is derived from the petE promoter from pea plastocyanin gene, by a deletion in % of the repeated invert sequences, of the as-1 like and enhancer like elements borne by two Sphl fragments of 403 bp. The petE promoter directs cell-specific but not full light-regulated expression in transgenic tebacco Flants. The promoter is used to construct chimeric promoters of the invention. The specification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; Page 67; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nucleic acid sequence
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                                                                                                Sequence 326 BP; 111 A; 91 C; 38 G; 86 T; 0 other;
                                                                                                                                         barley, bat, lide, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents the chimeric promoter Mpr1096.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         derived from a promoter of the pea plastocyanin gene
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                       Score 189 6;
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                                                       Arabidopsis thaliana DNA fragment SEQ ID NO: 7098
                                                                  17-OCT-2000
                                                                               AAC34566;
                                                                                          AAC34566 standard; DNA; 686
                                                                                                     7
                                                                  (first entry)
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Hybridisation assay; genetic mapping; gene expression control, protein identification; signal transduction pathway;

metabolic pathway; promoter; termination sequence;

Arabidopsis thaliana

06-SEP-2000

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                                                                                     identify compounds that inhibit or activate a polypeptide involved in induced resistance, by contacting the cell or polypeptide with one or more test compounds and identifying compound(s) that specifically modulate the activity of the polypeptide. Plant protection agents can be prepared by formulating the identified compounds in a form suitable for their uptake and optionally metabolization in organisms, especially plants, phytopathogenic lungi, nemalodes or insect pests The nucleic plants, phytopathogenic lungi, nemalodes or insect pests.
                                                                                                                                                                                                                         can be used to identify compounds useful DNA molecule containing a self with a recombinant DNA molecule containing a selectable and/or detectable marker linked to the regulatory region, culturing the cell in the presence of a test compound or sample that suppliesses or activates and/or enhances expression of the marker flost cells containing the nucleic acids, or polypeptides encoded by the nucleic acids, can be used to
              acids or regulatory regions, or corresponding vectors, or the polypeptides encoded by the nucleic acids, or compounds identified as above, can be used to improve the pathogen resistance of plants or to produce transgenic plants. The nucleic acids or regulatory regions, or
                                                                                                                                                                                                                                                                                                                                                                  This invention describes novel cDNA molecules corresponding to tobacco (Nicotiana tabacum) genes that are expressed in response to tobacco mosaic virus ('MY) infection. Regulatory regions that provide specific expression of the nucleic acids in connection with induced resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim la; 66; 130pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-552163/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schreier P, Garbers C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tobacco; plant; resistance; tobacco mosaic virus; TMV; infection; protection; plant protection agent, phytopathogenic fungi; rematode;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pathogen resistance in plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acids useful for identifying compounds capable of inducing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (FARB ) HAYER AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      insect pest; pathogen resistance; transgenic plant; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tobacco plant resistance-associated cDNA fragment 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ33777;
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cells, plants or plant tissues containing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 resistance in plants. Vectors containing the nacicic acids, or the regulatory regions of recombinant DNA molecule or vectors containing them, or the compounds identified as above, or the polypeptides or autibodies, can be used in diagnostic compositions. AAZ34076-233888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptides and antibodies to the polypeptides, can be used to screen for new plant protection agents or genes involved in induction of
                                                                                                                                                                            (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                             22-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      represent fragments of tobacco cDNA which are associated with plant
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                                                                                                                                           Gruber V,
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                                                                                                                                                                                                                                                                                                                                                                                           promoter; ss
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    Mismatches

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Pred. No. 3.2e-19;
Pred. No. 3.2e-79;
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Claim 2; Page 67; 83pp; English

derived from a promoter of the pea plastocyanin gene monocotyledonous transgenic plants comprises a nucleic acid Chimeric expression promoter for producing directyledonous

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The present sequence represents the chameric promoter Mprilips. The promoter is derived from the petE promoter from pea plastocyania gene,
                                                                                           Chimeric expression promoter to producing directyledomous and monocotyledomous transferice plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin dese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression promoter comprising a promoter of the pea plastocyanin generations comprising a 6-box operably or functionally linked upstream of a CANT box, TATA box and transcription intiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledomous species, e.g. potato, tobacco, conton, lettuce, tomato, meion, cummber, pea, rape, bestroat, or sunflower, and memocotyledomous species, e.g. wheat, battley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              promoter is derived from the petE promoter from ped plastocyanin gene, and contains only the INTA and cAAT boxes, and corresponds to a minimal reference promoter on the petE promoter. The petE promoter directs cell specific but not full light regulated expression in transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric
                                                          Claim 2: Page 69: 83pp: English
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                                                                                                                                                                              WP1: 2000 587667/55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        petE promoter; chimeric promoter; transgenic plant; MPrl109;
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RESULT 1
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Matches 138; Conservative
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                                                               Bensen RJ:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transit_peptide [..43%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Malze: plastid targetting sequence, transit pertide plastocyanin; subcellular localisation; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression vectors for producing transgenic plants, such as directlyledonous species, e.g. potato, tobacco, coston, lettuce, tomato, melon, cucumber, pea, rape, beetroot, or sunflower, and monocotyledonous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a promoter of the pea plastocyanin gene, or comprising a G box operably or functionally linked upstream of a CAAI box. [A]A box and
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WDZ000112732 AZ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zed mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Maize plastid targetting transit peptide DNA (7)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ51834 standard; DNA; 437 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transcription initiation site. The chimeric promoters are used in
                                                                                                                                   (PION ) PIONEER HE BRED INT INC
                                                                                                                                                                                                                  28 AUG 1998;
                                                                                                                                                                                                                                                                                              25 AUG 1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 250 BP; 81 A; 67 C; 31 G; 71 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     species, e.g. wheat, barley, pat, rice, or corn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   850 Babbar Conditionate that according to an analysis of the following section of the following section of the following section and according to the following section of 
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/transl_except (posijo..i5, aa:Xaa)
/note="Xaa Any amino acid"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cuery Match 7.4%;
Fest Local Similarity 65.6%;
        Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pisum
                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                               Nucleotide sequence of the promoter MPr1108
                                                                                                                                                                                                                                                                                                                                                       08-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 437 BP; 75 A; 156 C; 121 G; 69 T; 16 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             particular, the transit peptide finds use in the localisation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present DNA encodes a maize transit peptide which has homology to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 7, Page 44; 50pp: English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New organelle targeting nucleic acid and amino acid sequences are useful in genetic engineering for modulating the subcellular localization of heterologous proteins in plants.
                                                                                                                                            22-MAR·1999;
                                                                                                                                                                     20-MAR-2000; 2000WO-IB00317
                                                                                                                                                                                               28-SEP-2000
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                                                                                                                                                                                                                                                                                       plastocyanín gene promoter; ss
                                                                                                                                                                                                                                                                                                                                                                                                             AAA96461 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plastocyanin.
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                                                                                                                 (MERI-) MERISTEM THERAPEUTICS.
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                                                               2000-587667/55
                                                                                                                                                                                                                                                                                                   promoter; chimeric promoter; transgenic plant; MPr1108;
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                                                                                        Gruber V,
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                                                                                                                                            99FR-0003635
                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                          Theisen M;
                                                                                                                                                                                                                                                                                                                                                                                                              281 BP
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Pred. No. 8.3e-11;
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Best Local Similarity 77.8%;
Matches 133; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    promoters of the invention. The specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a Gox operably or functionally inted upstream of a CAAI box, TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, nelon, cucumber, pea, rape, bectroot, or sunflower, and monocotyledonous species, e.g. wheat, barley, oat, rice, or corn.
                                            Chimeric expression promoter for producing dicotyledonous and monocutyledonous transgenic plants comprises a nucleic acid siderived from a promoter of the pea plastocyanin gene \,
                                                                                                                                                             Rance I,
                                                                                                                                                                                                                                                                                                                                                                                                                                          plastocyanin
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                                                                                                                          WPI; 2000-587667/55
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                                                                                                                                                                                                                                                                          20-MAR-2000;
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                                                                                                                                                                                                  (MERI-) MERISTEM THERAPEUTICS
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                                                                                                                                                               Theisen M;
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U: Mismatches 36
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Claim 2; Fage 70; 83pp; English

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AAA96465
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Chimeric expression promoter for producing dicotyledonous and monocotyledonous transferic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin mene
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transgenic tobacco plants. The promoter is used to construct chimeric
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                                                                                                                                                                                                                                                                           WPT: 2000 587667/55
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                                 Query Match 6.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             promoter is derived from the petE promoter from pea plastocyanin dence, by lusing the petE as I like and nos enhancer like elements to the promoter MPr1000 (comprising TATA and CAAT boxes of petE) and then inserting a 6 box and lusing a fragment comprising a diplication of the element as2 and as1. The petE promoter directs cell specific but not clement as2 and as1.
                                                                                                                                                                                                                                                                                                                           specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin deno, or comprising a 6-box operably or functionally linked upstream of a CACT box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                   tal. Light regulated expression is transposed totally plants. The promoter is used to construct chimeric promoters of the insention. The
                                                                                                                                                                                                                                                        tor producing transferde plants, such as dicoryledonous species, e.g. polato, tobacco, cotton, lettuce, tomato, melon, engumber, pol
                                                                                                                                                      Sequence 296 BP: 94 A: 74 C: 45 G: 83 T: 0 other:
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## SUMMARIES

Fesult No.	Seute	Query Match	Query Match Length DB	В	[I]	beset iption
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13	375.4	27.8	553	10	BE317514	BE317514 NF040F10L
14	375.4	27.8	555	9	AW683640	AW633640 NF017A04L
15	375.4	27 A	53.5	10	BF637258	BF537058 NFC715031
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## ALIGNMENTS

	Source	FEATURES							COMMENT	JOURNAL		TITLE	AUTHORS	REFERENCE				ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	OFFINITION	FOCUS	BI271179	RESULT 1
/organism= medicago truncatura /db_ripf="tayon:380" /clone="NFO50C10FL" /clone_lib="Developing llower" /tissue_type="Developing llowers" /de:_staye="Teveloping llowers" /de:_staye="Teveloping, faily opened flowers and flowers in early transition into pods."		SE [FI DOT I W W WENAW MS INTENT.] Location/Qualitiers 1	Plate: 050 row: C column: 10	Email: gdmay@noble.org	Fax: 580 221 7380	IDIO SAM MODIO FARRWay, Archote, CR (34CD) CSA Tel: 520 DDI 7391	le Foundation	Plant Biology Division	Contact: May GD	Unpublished (2001)	Medicago truncatula flower library	Expressed Sequence Tags from the Samuel Roberts Noble Foundation	Torres Jerez, I. Scott, A.D. Harris, A.R., Gonzales, R.A., Bell, C.J.,	1 (bases 1 to 611)	Medicago.	Spermanophyna; Madhollophyna; endiconyledons; core endicons; Posidae: enrosids I: Fabales: Fabaceae: Papillopoideae: Tritolipae:	Rukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;	Medicago truncatula	barrel medic.	EST.	BI271179:1 GI:14879432	BI271179	NEUSUCIUELLEIUBZ Developing flower Medicago truncatula cDNA clone			

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DEFINITION
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                                                                                                           1086 tgccaaggctttgtgtcagagcttccctcaaagactttggagttqctqctqttgccactq 1145
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The Samuel Roberts Noble Foundation
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Seq primer: TCACACAGGAAACAGCTATGAC
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Fax: 580 221 7380
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/note="vector: Lambda Zap; At the trifoliate stage, M.
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The Samuel Roberts Noble Foundation
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Ouery Match 2008: Seare (97.5) Let 16, Length 645; Hest Local Similarity 96 18; Pred. No. 16 55; Matches 421; Conservative 9; Mismatches 9; Indels 8;
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Tel: 580-221 [ci]
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                                                                                 Plant Biology Division
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                                                                                                                                                                                                                                                                 Rosidae; eurosids 1, Fabales, Fabadeae; Papilionoideae, Trifolieae
                                                                                                                                                                                                                                                                                                                                      barrel medic
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              Tel: 580 221 7302
Fax: 580 221 7380
                                                                                                                                                                 Expressed Sequence Tags from the Samuel Roberts Noble Foundation
                                                                                                                                                                                              (bases 1 to 603)
X -Z , Shadle,G , Scott,A D , Harris,A P , Gonzales,P.A , Bell
J , Flores,H P , Inman,J.T., Weller,T W , May,G D and Dixon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
radixon@noble.org
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/note="Vector: Lambda Zap; Contains a mixture of very
young, developing, mature and senescing leaves."
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                                                                                                              2000 this sequence version
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Fedorova,M., Pierson,B.L., Samac,D.A., Vance,C.P., Gantt,G.S.,H., Ellis,L., Town,C.D., Bownab,C.L., Craven,M.B., Hansen,L.S. Holt,I.E. and Fraser,C.M.
ESTs from leaves of Medicago truncatula after inoculation with
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/foole="Vector Lambda Zap, Contains a mixture of internodal stem segments"
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/tissue_type="stem"
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Pred. No. 3.5e-54;
Mismatches 8;
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Fax: 651 649 505B
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Modicago Genome Initiative accession: McL:S:14935
Insert Tempth: 7-8 Std Error: 0.00
Plate: 016 row: B column: 04
Seq primer: TYMCAMAMAMAMATATATAM.
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The Samuel Roberts Noble Foundation
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161: 580-221-7312
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//clone_lib-"Developing rood"
//lissue_type="root"
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/lost "Wether loaded Ear Load EBA was extracted from non-inducted roots of plants grown in Lamb citrate medium. Samples were taken at four time points (approximately two days, one, two and six weeks post growing from prepresenting early seedling growth to bitrogen_limitation."
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                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                     NP053R06LP1P1045 Developing leaf Medicago trumatula cDNA clone
NP053B06LP 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: gdmay@noble.org
Mcdicago Genome Initiative accession: MGI:S:21826
Insert Length: 653 Std Error: 0.00
Plate: 053 row: B column: 06
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Tel. 580 221 7381
Fax: 580 221 7380
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The Samuel Roberts Noble Foundation
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Flores,H.R., Inman,J.T., Weller,J.W. and May,G.D.
Expressed Sequence Tags from the Samuel Roberts Noble Foundation
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Spermatophyta, Magnoliophyta, cudicotyledons; core eudicots;
Rosidae; eurosids I; Fabales; Fabareae; Papilionoideae; Trifolieae
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/tissue_type-"leaf"
/tissue_type-"leaf"
/dev_stage="Pooled developmental"
/note "Vector. Lambda Zap, Contains a mixture of very
young, developing, mature and senescing leaves."
a 108 c 106 q 135 t
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NEO67H08LF 5', mkNA sequence.
BE316878
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Medicago Genome Initiative accession: MGI+S-1895]
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    Expressed Sequence lass from the Samuel Roberts Noble Foundation

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2510 Sam Noble Parkway, Ardmore, OK 73402,
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Eukaryota, Virioiplantae, Streptophyta; Embryophyta; Iracheophyta;
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                                                                                                                                                                        Email: qdmay@noble.ord
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The Samuel Roberts Noble Foundation
2510 Sam Noble Parkway, Ardmore, OK 73402, USA
Tel: 580 221 7391
Fax: 580 221 7380
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/clone_lib="Developing leaf"
/tissue_type-"leat"
/dev_stage="Pooled developmental"
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/clone="NF017A041F"
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                                                                                                                                                                                                                                                                                                                                                                                          Medicago truncatula
Eskaryota: Viridiplantae: Streptopbyta: Embryophyta: Tracheophyta:
Spenmatophyta, Magnoliophyta, codicotyledona, core endicots:
Rosidae: eurosids I: Fabales: Fabancae: Papilionoideae: Fritolieae;
                                                                                                                                                                                                                                               Unpublished (2000)
Contact: May GD
                                                                                                                                                                                                                                                                                                 Torros-Toros,I., Scott,A.D., Harris,A.R., Genzales.R.A., Bell.C.J. Elores,H.R., Isman,J.T., Weller,J.W. and May,G.D. Expressed Sequence Tags from the Samuel Roberts Noble Foundation
                                                                    Email: gdmay@noble.orq
Insert Length: 566 Std Error: 0.
Plate: 072 row: 6 column: 03
Seg primer: TCACACAGGAAACAGCTAICAC
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The Samuel Roberts Noble Foundation
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                                                                                                                                                   Tel: 580 221 7391
Fax: 580 221 7380
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/Grqanism-"Medicago trubcătulă"
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NUMBER OF SEQUENCES: 45  CORRESPONDENCE ADDRESSE: Knobbe Martens Olson & Bear CITY: Newport Beach COUNTRY: US ZIP: 92660  STATE: California COUNTRY: US ZIP: P92600  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: LIM FOR COMPALIA: COMPUTER: Floppy disk COMPUTER: OBSECTION DATA: P0-105/MS-105 SOFTWARE: ISTECTION DATA: P0-105/MS-105 VCTSIGN #1.  CURHENT APPLICATION DATA: ADPLICATION: 435  PILING DATE: 10-SEP-1993  CLASSIFICATION NUMBER: US/85/8487.826B  FILING DATE: NUMBER: NIHIZI.001CP1  TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION STORMS 13: GEOGRAPHE COLORS 13: GEOGRAPHE COLORS 13:	UESULT 1  S-08-487-826H-13  Sequence 13, Application Patent No. 5993827  GENERAL INFORMATION: APPLICANT: Sim, Kim I APPLICANT: Chitchis, G APPLICANT: Miller, Ld APPLICANT: Su, Xim I APPLICANT: Su, Xin I APPLICANT: Su, Xin I TITLE OF INVENTION: F	28
	SNIGLONG ONIGHTA GLADOMENTONIS XVALA MOTOOMEN	Sequence 5, Appli Sequence 11, Appl Paten: No. 5231168 Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 1, Appli Sequence 1, Appli Sequence 3, Appli Sequence 3, Appli Sequence 5, Appli Sequence 2, Appli Patent No. 5175495 Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli

HYPOTHELICAL: NO CONTROL NO CONTR

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SEQUENCE CHARACTERISTICS:
LENGTH, 19124 Luse pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                             Sequence 14. Application 05/08, 32463 Patent No. 5670367
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                                                                                       GENERAL INFORMATION:
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              APPLICANT: SCHELELLINGLE APPLICANT: FALKNER, F G
  TITLE OF INVENTION:
                                                                 APPLICANT:
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RECOMBINANT POWLECK VIRUS
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APPLICATION NUMBER: US/07/935, 413
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OPERATING SYSTEM: PC-bos/MS bos
SOUTWARE: Patentle Release #1.0, Version #1.26
CORRENT APPLICATION DATA:
1441 KERKHRHERHERRKKKREPREPREPRERREPRERREPRESERENDERKERENDERER (1292
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ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
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                                                                                      NUMBER OF SECHENCES:
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                                                                                                                                                                                                                                        Query Match 4.9%; Score 65.8; DB 4; Length 837; Best Local Similarity 46.7%, Pred No. 5.66.07; Matches 279; Conservative 0; Mismatches 312; Indels
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                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
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LENGTH: 837 base pairs
TYPE: nucleic acid
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
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NAME: Meigs, J. Timothy
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APPLICANT: Knechtle, Fhilip
APPLICANT: Rebischung, Corinne
TITLE OF INVENTION: GENUMIC INA SEQUENCES OF ASHBYA GUSSYPII
TITLE OF INVENTION: AND USES THEREOF
NUMBER OF SEQUENCES: 1152
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PRIOR APPLICATION DATA:
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188 thttgttgcaacatttgagaaaattttgttgttctctcttttcattggtcaasaacaata 247
                                                                                                                   128 aaacattagagtaaagaaatatggatgataagaacaagagtagtgatattttgacaacaa 187
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DUS/MS-DUS
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CITY: Research Triangle Park
STATE: No. 6239264th Carolina
                                                                               68 aagtgtgtacatttttatttgaacaaaaatattcacctactactyttataaatcattatt 127
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                                                                                                                                                                                                                                                                                                                                                                       ORGANISM:
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Steiner, Sabine
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                   INFORMATION FOR SEQ ID NO:
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APPLICANT:
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                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                     COMPUTER: TEM PC COMPATIBLE
OPERATING SYSTEM. PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.40
CURRENT APPLICATION DATA
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                                                                                                                                                          FILING DATE: 31-DEC-1996
ATTORNEY/AGENT INFORMATION:
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SEQUENCE CHARACTERISTICS:
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                                                                                               NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: PF
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Steiner, Sabine
Mohr, Christine
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Knechtle, Philipp
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                                         CORRESPONDENCE ADDRESS
                                                                                    TITLE OF INVENTION: OF NOMITED ASSOCIATIONS OF ASHRYA GOSSYPTI
                                                                  NUMBER OF SEQUENCES:
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TYPE: medeic acid
STRANDEDNESS: Single
                      ADDRESSEE:
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3054 Fornwallis Foad
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Knechtle, Flalig
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                  No. 6239243ar s Corporation
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TELECOMMORICATION INFORMATION
TELEPHONE: 919-541 8587
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496 ttaaacatgtqattaIIItaalqaaftqatqaaagaqttqqaftaaauffqtattaqtaa 594 i.e. i.e. i.e. i.e. i.e. i
                                                                    436 tigaraayteattaaaaaaaaaaaaattattttaaaattaaaattaaattaaattaaattaaa
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                                                                                                                                                                                                                                                          256. AATAATATETIAATAACIAA HIJAAAA UTGAACA JAGACIAAA IACIA HEAHAAA HEE
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OPERATING SYSTEM: PC DOS/MS DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: CH 0016/97
FILING DATE: 31 DEC 1996
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STAIR: No. 6239264th Carplina
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER
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EDNESS: single
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US-08-998-416-1137
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                                                                                                                                                                                                                                                                                                    Matches 245;
                                                                                                                                                                                                                                                                                                                                          Query Match
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LENGTH: 636 base pairs
TYPE: nucleic acid
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 919-541-8587
TELEFAX: 919-541-8689
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
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                                          128 aaacattagagtaaagaaatatgyatgataagaacaagagtagtgatattttgacaacaa 187
248 gagagaaaaaaggaagaggagaataaaaacataatgtgagtatgagagaaagutgt 307
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                                                                                    188 ttttgttgcaacatttgagaaaattttgttgttctctcttttcattggtcaaaaacaata 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                    68 aagtgtgtacatttttatttgaacaaaaatattcacctactactgifataaatcattatt 127
                                                                                                                                                                                                                                                                                              / Match 4.4%; Score 58.8; DB 4; Length 636;
Local Similarity 45.5%; Pred. No. 2.4e-05;
hes 245; Conservative 0; Mismatches 292, indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: PF/5-30306/A/CGC1976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: CH 0016/97 FILING DATE: 31-DEC-1996
                                                                                                                            STRANDEDNESS
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Steiner, Sapine
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Knechtle, Philipp
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FEMTION: GENOMIC DNA SEQUENCES OF ASHBYA COSSYPTI
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N: 435
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        Matches
                      Query Match
Best Local Similarity
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                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 8920 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.24 CURRENT APPLICATION DATA:
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APPLICANT: O'Sullivan, William J
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                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 485 TEFCTTATTATTAATTTTTATATTATTTAATTAATTATATTCATTTTATTTATTTATT 542
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REGISTRATION NUMBER: 29.009
REFERENCE/DOCKET NUMBER: 47-80
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                    TOPOLOGY:
                                                                                                                                                    STRANDEDNESS:
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TELEPHONE: 703-816-4100
                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US, 0 FILING DATE: 06-Jul-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Arlington
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    226,
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1100 No. 5849573th Glebe Road, 8th Floor
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    Conservative
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Score 57.2; DH 2;
Pred No 0.00015;
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EARLIER APPLICATION NUMBER: 15-6-380
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                                                               INFORMATION FOR SEQ ID NO: 14: SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: AND PLASMODIUM PARTITIONARM GEVERN VIVAX
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      TITLE OF INVENTION:
                        APPLICANT:
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ANTI SENSE. NO
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                                                                                              Pohlmann, Rainer
Steiner, Sabine
                                                                                                                                       Philippsen,
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                                                                            Mohr, Christine
                      Rebischung, Corinne
                                        Wendland, Jurgen
Knechtle, Philipp
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GENOMIC DNA SEQUENCES OF ASHBYA GOSSYPII
                                                                                                                                       Peter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 3.9%; Score 53.2; DB 4; Length 6
Best Local Similarity 47.7%; Pred. No. 0.00052;
Matches 251; Conservative 0; Mismatches 260; Indels
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INFORMATION FOR SEQ ID NO: 1137:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 31-DEC-1996
ATTORNEY/AGENT INFORMATION:
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449 aaaagaaagaataaattattttaaaattaaaagttyagtratttgattaaacatgtgat 508
                                      483 TAAAGTTAAATTTAATAATAATAATTCTTATAAAAAAGATTAAATAATATAAATCAAC 424
                                                                                                                 243 TAACTAATTTAAAATTTGAACATAGACTAAATAGTATTCATATTAAATATTATTTTATA 184
                                                                                                                                                                                            152 algataagaacaagagtagtgatattttgacaacaattttgttgccaacatttgaqaaaat 211
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                                                                         389 aataaataaggatgacgcattagagagatgtacmattagagagattttliggcaagtcatta 448
                                                                                                                                                                                                                                                                       TELEPHONE: 919-541-8689
                                                                                                                                                                                                                                                                                                                                                                                                                                                              94 aaatattcacct--actactgttataaatcattattaaacattagagtaaagaaatatgg 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         34 aaattaactoaaaaaccataaaagtttaagttagcaaqtqtqtacatttttatttqaacaa 93
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 38,241 REFERENCE/DOCKET NUMBER: PF
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OPERATING SYSTEM: PC-DOS/MS-DOS
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Research Triangle Park
No. 6239264th Carolina
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 Best Local Similarity
                Query Match
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                                                                                                                  FEATURE:
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APPLICATION NUMBER:
FILING DATE: 30 JAN
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SUFTWARE: Parentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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ADDRESSEE: David R. Saliwanchik
STREET: 2421 N.W. 41st Street, Suite A l
                                                                                                                                                                     FEATURE
                                                                                                                                                                                                                       FEATURE
                                                                                                                                                                                                                                                       MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
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REGISTRATION NUMBER: 41, 3
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EDNESS: double
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 5721352
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                     APPLICANT: Gruidt, Michael E.
TITLE OF INVENTION: No. 9721952el Entomognoxyrias Expression System
NUMBER OF SEQUENCES: 40
                                                                                                 COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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                   MUDIUM TYPE: Floppy disk COMPUTER: THM PC COMPATIBLE ASSESSED.
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Y: U.S.A.
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242] N.W. 41St Street, Synte A 1
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PatentIn Release #1.0, Version #1.25
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APPLICATION NUMBER: US 07/827,658
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NAME/KEY:
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1035 AAAAGTTTATATATTTCAAATTTAAATATTATTAATTTAAATTTATAACAAAAATTAAAA 1094
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                                     462 aattatttttaaaattaaaagttgagtcatttgattaaacatgtgattatttaatgaatt 521
                                                                            975 GAAGAATTGCATATAATATTATGATAATAATATTTTTAAATAATATTCCAGAAAATATT 1034
                                                                                                                                                           915 ATAATAATAAACAGAGAFAATGTTATTAATATTAATATTAAAAAAAATTAGTTAATTTA 974
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                                                                                                               342 attgaggaatttgacaasagctacacaaataagggttaattgctgtsaafaaatsaggaf 401
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Local Similarity 44.7%; Pred. No. 0.0014;
hes 298; Conservative 0; Mismatches 355; Indels 13; Caps
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REFERENCE/DOCKET NUMBER: UF114.C2
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REGISTRATION NUMBER: 31,79
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Patent No. 5935777
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                                                                                           INFORMATION FOR SEQ ID NO: 8 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 904-375-8100
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APPLICATION NUMBER: 1
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MOLECULE TYPE:
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                   TOPOLOGY: uni
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2421 N.W. 41st Street, Suite A-I
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RESULT 14
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                                                 Sequence 288, Application 13/28/08/08416
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                                                                                                                                                                                                                                                                                                     1211 ΤΑΙΤΑΑΤΑΑΤΗ ΑΘΙΑΑΑΣΗ ΑΛΛΑΑΑΑΑΙ ΙΑΑΙΑΑΙΑΤΑΤΑΑΑΛΑΑ ΛΑΑΤΙΤΙΘΙΑΑΘΤΙ 1270
                                                                                                                                                                                                                                                                                                                                                                                                      1155 CONCATTOTATAGAATTI FIAAA FEGIGAATOATGIAA FATAAATGACTATAATTI -
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                                                                                                                                                                                                                                                                                                                                                  642 tagatcaaalaagagaaalaacgalalattaatcectecsaaaaaaaaaaaaaggtatatt 201
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LOCATION:
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Philippson, Peter
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Prod. No. 0.0014:
0: Mismatches 565, Indels 13: Gaps
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NAME: MOTORS NOMBER: 38,241
NECLES RATION NOMBER: 38,241
HELES MANINICATION INFORMATION:
TELES MANINICATION INFORMATION:
TELES MANINICATION INFORMATION:
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APPLICANT: Knechtle, Pullipp
APPLICANT: Rebischung, Corline
(1913) OF INVENTION, JEROMIC DNA SEGURATES DE ASHRYA DESSYPT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: HAM PC compartable
OPERATING SYSTEM: PC DOS/MS DOS
SOFTWARE: Parenth Release #1.0, Version #1.co
CURRENT APPLICATION DATA:
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173 ataitittyaeaaeaatiittyittyeaaeairttyayaaaatiittyifyttetetetiitteat 232
                                                                                                                                                                                                                                                                                                                                                                                  462 TAATAATTCTTATAAAAAAGATTAAATAATAAAATCAACATAATATTATAAAAAATACAT 407
                                                                                                                                                                                                                                                                                                                                                                                                                      113 ttataaatraltattaaarattägagtaaagaaatatggatgataagaacaagaagtagtg 122
                                                                                                282 APTHAPTAAAGAAAATAATAATAATAATAATAATATTITAATAACTAATTIAAAAATTICAAC 22 -
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CHY: Research Triangle Park
SIAIE: No. 6239264th Carolina
                                                                                                                                                                                                                                                                                                                                                                                                                                                            y Match 3.8%; Score 51.8; D8 4. Legath 80%; Local Similarity 47.9%; Pred. No. 0.0012; hdc/s 210; Conservative 0; Mismatches 220; Hdc/s 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
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Steiner, Sabine
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US-38-998-416-595/c
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                                                                                                                (uery Match 3.6%; Score 48.8; HB 4; Length 558;
Eest Local Similarity 47.3%; Pred. No. 0.0059;
Matches 210; Conservative 0; Mismatches 232; Indels
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APPLICANT:
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APPLICANT: Knechtle, Philipp
APPLICANT: Rebischung, Coritine
TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHBYA GOSSYPII
TITLE OF INVENTION: AND USES THEREOF
NUMBER OF SEQUENCES 1152
                                                                                                                                                                                                                                                      MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: PT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
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PRIOR APPLICATION DATA:
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150 ggatgataagaacaagactagtgatattttgacaacaattttgttgcaacatttgagaaa 209
                                         530 agttggattaaagttgtaitagtaa 554
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                                                                                                                                                                                                                                                                                                           LENGTH: 658 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: CH 0016/97 FILING DATE: 31-DEC-1996
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FILING DATE: 24-DE
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Steiner, Sabine
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                                                                                          aaaagaaagaataaattattttaaaattaaaagttgagtcatttgattaaacatgtgat
                                                                                                                                                         ataaataa-ggatgacgcattagagagatgtaccattagagaattttttggcaaqtcatta 448
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Search completed: September 1, 2002, 00:57:11 Job time, 7903 sec

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 20000ວິດຄວິດ
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Perfect score:
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                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Query
Score Match Length DB ID
                                                                                                        1797656 seqs, 10463268293 residues
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## ALIGNMENTS

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BASE COUNT ORIGIN	FEATURES Source	REFERENCE AUTHORS TITLE JOURNAL	VERSION KEYWORDS SOURCE ORGANISM	AX105268 TOTIS DEFINITION ACCESSION
<pre>/db_xref="taxon:32630" /note="Sequence to be used as a Promoter for requiating expression" 390 a 141 c 145 q 295 t</pre>	medicago inc. (cA) Location/Qualifiers 1971 /organism-"synthetic construct"	1 (bases 1 to 971)  Vezina,L.P. and D'Aoust,M.A.  Promoter for regulating expression of foreign genes  Patent: WO 0125455 A 2 12-APR-2001:	AXI05268.1 GI:13921418  synthetic construct. synthetic construct artificial sequence.	AX105268 471 bp NA Sequence 2 from Patent W00125455 AX105268
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vezina, I P and D'Arust, M A
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/note="Sequence to be used as a expression"
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Rance.I., Theisen.M. and Gruber,V.
Patent: FR 279388-A 2 29-SEF-2000;
MERISTEM THERAPEUTICS (FK)
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1 (bases 1 to 426).
Rance, J. Theises, M. and Scuber, V.
Patent: FR 2791458 A 4 29 SEP 2000;
                                                                                                                                                                                                                                                                                   Submitted (09 SEP 1992) J. Gray, University of Cambridge, Dept. of Plant Sciences, Downing Street, Cambridge CK, SEA, UK
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/prote-n_id="CAA48393.1"
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25, .62
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813. .1319
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134. .171
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/clone_lib="lambda EMBL3"
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/clone-"lambda-GPC3"
/product="mature plastocyanin (AA 1-99)"
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/MORE-The region is covered only by PCR products which were amplified with a high fidelity polymerase and seguenced with both dye terminator and STP terminator
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DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              on Jul 13, 2001 this sequence version replaced di:14578148. This sequence was finished as follows unless otherwise noted: all regions were either double stranded or sequenced with an alternate chemistry or covered by high quality data (i.e. phred quality >-30), an attempt was made to resolve all sequencing problems, such as compressions and repeats, all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the assembly was confirmed by restriction digest.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Huang, R., de la Hastide, M., Spiegel.i., Nascimento, I., Kirchoff, K
Preston, R., King, I., Zutavern, I., Santos, I., Miller, B., Kuit, K.,
Cunnius, D.M., Bell, M., Halija, V., Shah, R., Hahret, A., Yang, C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (30 JUN 2001) Lita Armemberg Hazer Genome Sequencing Center, Cold Spring Harbor Laboratory, 1 Bungtown Road, Cold Spring Harbor, NY 11724, USA
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Preston,F., King,L., Extareco,T., Santos,L., Milley, E., Kuff,K.,
Cunnius,D.M., Bell,M., Halija,V., Shah,R., Habret,A., Yang,C.,
Palmer,L., O'Shaughnessy,A., Dedhia,N. and McCombie,W.R.
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Submitted (13-301-2001) Lita Annenberg Hazen Genome Center, Cold
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Spring Harbor Laboratory, 1 Bungtown Rd, Cold Spring Harbor, NY
/note-"The sequence is covered by multiple subclosed derived from a PCR product amplitted with a high tidelity polymerase. The sequence is of high quality."
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/db_xref="taxon:3880"
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Hest Local Similarity 69.2% Fred. No. 1.6e 08;
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Sequence 6 from Patent FR2791458
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                                                                                                                                                                   1 (bases 1 to 280)
Rance, L. Thorsen, M. and Gruber, V.
Patent, FR 2794 (88-A 7 29 SEF 2000)
MDRISTER THERAPEUTICS (FR)
                                                                                                                                                                                                                                                                                                                                           Sequence / from Patent FR2791358
AXU36603
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a 'G' box and by fusing a sequence of 44 bp of the RNA. \operatorname{\mathsf{CS}} came promoter
                                   position 99 bp of MPT1098 an element of 18 bp containing
                                                                                        /ordanism "synthetic constract"
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Search completed: September 1, 2002, 00:55:18 Job time: 17704 sec

GenCore version 4.5 Copyright (c) 1993 - 2000 Compages Ltd

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SUMMAPIES

Result	)	Query				
N <sub>C</sub> .	Score	Match	Match Length DB		ID	Description
	971	100.0	971	()	AAS02127	Promoter #2 for re
ıى	971	100 n	058.1	12	MS02126	Fromoter #1 tor re
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4	191.8	19.8	<b>6</b> 13	: 1	AMA9645B	Nucleotide sequen
υī	191.8	19.8	8:4	21	AAA96457	Nucleotide sequenc
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Human immune byote	ABL33671	24	8445	6.3	ଞ୍ଚ. ଜୁ	4 л	ว	
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Human immune syste	ABL33943	24	12138	6.3	61.2	43		
Homan immune syste	AHL33994	24	6483	6.3	61.4	42		
Tumour suppressor	AAS46595	.)	5445	E)	61 4	4 1	2	
Human gene regulat	AAS61077	24	5019	у. Д	۶ <u>1</u> 8	<b>4</b>	3	
Human immune syste	ABL32326	1.0 #¥	6103		51.8	3	7.	
"Cumour suppressor	AAS46717	64 64	19131	ক ্ৰ	62.8	38		
(umour suppressor	AAS46746	F3	38342		20.00	37		
Human gene regulat	AAS61128	24	7913	6.5	63	36		
Tumbur happressor	AAS46311	ta ta	7913	5.5	<u>ش</u> ري	ш :л		
	ABL34025	24	6375	6.5	63	34		
Muman immune syste	ABL33294	24	17419	5.5	63.6	w w	<b>a</b>	
	AAS45392	t. t.	27419	ir	t 4.5	ŭ L	IJ	
	ABI.32028	<u>د</u>	7351	6.5	63.6	ω, 	C	
Human immune syste	ABL32972	ن 4	(* : 4 F	(* , #	53.5	30	c.	
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Tumber suppressor	AAS46429	ro N	6196	4.5	£3.8	27	C	
Human immune syste	ABL32098	<u>د</u>	1966	6. 7	65.2	56	c	
	ABI.33307	1) 4	6175	6.8	66.4	N 5	C	
Human immunc/hacma	AAK? 3165	tu tu	20420	7.0	68	۱.) 4 <u>.</u>		
Flusmodium var-7 p	AAZ 98287	P	19124	7.0	F8.2	ta W		
Plasmodium war 7 q	AATTIREI	3	19124	7.0	f.80.12	13		
Tumbur Suppressor	AAS46735	t2 t2	6292	7.0	58.2	11	0	
Human immune syste	ABI.32267	ده 4	6419	7.	68.8	20	C	
	AAK73166	te te	19465	1.4	) 1 . B	ī		
Human immune syste	AH1.34 358	() 4	12237	7.4	72	<del></del>	G	
Himan immuse syste	VBT33613	(1 «h	7597	7.7	74.4	17	3	
Mucleotide sequenc	AAA96477	F-1	259	7.8	76	16		
pNFX 0 xylanase cD	AAQ53480	15	2503	8.0	77.2	15		
l'isum sativum plas	AA1:77099	18	266	œ ы	79.2	4		
	AAA96466	-	220	9.1	88.4	13		
₩agleetide sequenc	AAA9£4£4	د.	202	£	28	12		
Mucléotide sequenc	AAA96465	i)	296	9.2	68	11		
#.cleotide sequenc	WW36463	1	1. 30 0	9.1.	8.9	10		

## ALTGNMENTS

RESULT 1 AAS02127

AAS02127 standard; DNA; 971 BP

18-JUL-2001 AAS02127;

(first entry)

Novel promoter for regulating expression of foreign genes organisms, more specifically in a leaf specific manner in WPI, 2001:266316/27. 04-00T-1999; Promoter #2 for regulating expression of foreign genes Vestina I, (MEDI-) MEDICAGO INC. 02-909-2000; 2000WO-CA01144. 12-APR-2001. W0200125455-A2 Synthetic Fromoter, transpenie plant, sonocetyledon; dicetyledon; gymnosperm; ds E'Adust M; 99115-7129 in transgement

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    Matches 241; Conservative
                  Query Match (1988)
Best Local Similarity 82.8%;
                                                                                                                                                                        expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operately or functionally linked upstream of a CAAF box. TATA box and transcription initiation site. The chimeric promoters are used in expression sectors to producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, softon, lettuce, tomato, melon, cucumber, pea, rape, bectroot, or suntlower, and monocotyledonous
                                                                                                                                                                                                                                                                                                                                                      promoter is derived from the petE promoter from pea plastocyanin dene, by a deletion in 5% of the repeated invert sequences as well as the as I like hox borne on a Spb. fragment of 212 bp. The petE promoter directs cell specific but not tall light regulated expression in
                                                                                                                                                                                                                                                                                                                 transgenie tobacco plants. The promoter is used to construct chimeric promoters of the invention, the specification describes a chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; Page 66; 84pp; English
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                                                                                                                Sequence 623 BP; 230 A; 128
                                                                                                                                                          species, e.g. wheat, barley
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                                    expression vectors for producting transposic plants, such as dicotyledonous species, e.g. potato, tobacco, conton, lettuce tomato, melon, cucumber, pea, raja, beetroot, or sunflower, and mongoutyledonous
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plastocyanin dene. The promoter directs cell-specific but not tull
light regulated expression in transferic tobacco plants. The promoter
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species, e.g. wheat, barley, oat, rice, or corn
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Query Match
Best Local Similarity
19.8%;
Score 191.8: I
Pred. No. 1.5e
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Sequence 834 BP: 289 A; [75 0] 133 G; 237 I; p other;

DR 217 (Sengts 834) Se 257

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RESULT
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beetroot, or sunflower, and monocotyledonous species, e g.
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   minimal reference promoter on the petE promoter. The petE promoter directs cell-specific but not full light-regulated expression in Figure 20.00 and the construct chimeric consequence of the permutet is used to construct chimeric
                                                The present sequence represents the chimeric promoter Mpr1098. The promoter is derived from the petF promoter from pea plastocyaning and contains only the TATA and CAAT boxes, and corresponds to a
                                                                                                          Claim 2; Page 67; 83pp; English.
                                                                                                                                     chimeric expression promoter for producing disotyledonous monosotyledonous transgenic plants comprises a nucleic ac derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                   WP1; 2000-587667/55
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The present sequence represents the chimeric promoter Mprilos. The promoter is derived from the potE promoter from per plastocyanin gene, by fusing the petE as 1 like and nos enhance—like elements to the promoter MPri098 (comprising IAIA and CAAI boxes of petE), and then deleting 34 be from the 57 UER. The petE promoter directs cell specific but not full light regulated expression in transpense tobacco plants. The promoter is used to expression in transpense tobacco plants. The promoter is used to expression in transpense tobacco plants in the specification describes a chimeron expression promoter comprising a promoter of the per plass.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2: Page 69; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chimerie expression promotes for producing disotyledonous and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   monocotyledomous transgence plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
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Prof. No. 46 12:
O: Mismatches
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expression promoter comprising a promoter of the pea plastocyanin ages, or comprising a 5 box operably of functionally cinked gistream of a 2AA box. IANA box and transcription initiation site. The chameric promoters
                                                                                      promoter directs cell specific but not full light regulated expression in
transienic tobacco plants. The promoter is used to construct climeric
promoters of the invention. The specification describes a chineric
                                                                                                                                                                                                                                                                                                                                                                                          Chimeris expression prometer to producing discryicotons and monocotyledomous transquair plants comprises a nucleic acid sequence derived from a promoter of the pea plastoryanth deno
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                                                                                                                                                                                promoter MPr1098 (comprising TATA and CAAL boxes of pert). The pert
                                                                                                                                                                                                                promoter is derived from the petE promoter from pea plastocyanin again
by fusing the petE as I like and new enhances like elements to the
                                                                                                                                                                                                                                                                          The present sequence represents the chimeric promoter Mp:1108. The
                                                                                                                                                                                                                                                                                                                                      Claim 2: Page 68: 83pp: English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rance 1, Gruber V,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleotide sequence of the promoter MFT1108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            species, e.g. wheat, barley, out, nice, or corn.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MERI ) MERISTEM THERAPEUTICS
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RESULT 1
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Best Local
                                                                 The present sequence represents the chimeric promoter Mpr1110. The promoter is derived from the petE juminter from pea plastocyanin gene, by fusing the petE as-1 like and nos enhancer like elements to the promoter MPr1098 (comprising 1ATA and CAAT boxes of petE), and then inserting a 6 box and fusing a fragmentof 35s RNA of CMV. The petE promoter directs cell-specific but not full light-regulated expression transgenic lobacco plan's. The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric
expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operably or functionally linked upstream of a CAAT box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as
                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                   Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nuclear acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, beetroot, or sunflower, and monocotyledonous
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                                                                                                                                                                                                                                   2; Page 70; 83pp; English

    Gruber V,

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77.88;
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                                                                                                                                    Chimeric expression promoter for producing dicotyledonous an monocotyledonous transgenic plants comprises a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        distilleduncias species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, beetroot, or sunllower, and monocotyledonous
                                                                                                                                                                        WF1; 2000-587667/55
                                                                                                                                                                                            Rance
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                                                                                                                                                                                                                                                                                                                                                                   petE promoter; chimeric promoter; transgenic plant; MPrllll;
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                                                                                                                                                                                                                                                                                                                                                          promoter;
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76.28;
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The present sequence represents the chimeric promoter Mprllll. The promoter is derived from the petE promoter from the plastocyanin gene, by fusing the petE as-1 like and see submarcer like elements to the promoter Mprl998 (comprising TATA and CAAI boxes of petE), and then inserting a G box and fusing a fragment comprising a duplication of the element as2 and as1. The petE promoter directs cell-specific but not full light-contact.

Claim 2, Page 71, 83pp, English.

derived from a promoter of the pea plastocyanin gene

full light-regulated expression in transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The

a chimeric expression promoter comprising

specification describes

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match (c.2%:
Best Local Similarity 76.2%:
Matches 144; Conservative
The present sequence represents the chimerre promoter Mprll53. The promoter is derived true the path promoter from pea plastocyania graditie present promoter directs cell specific but not full light regulated expression in Linspenie toleres plants. The promoter is used to construct chimerre promoters of the invention. The specification
                                                                                                                                                  Chimeric expression promotes for producing disolytedomous and monocotyledomous transfers e-plants comprises a nucleic acid sequence derived from a promoter of the pea plasforyaits dome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       initiation site. The chimeric promoters are used in expression vectors for producing transpense plants, such as dicotyledomous species, e.g. polato, tobacco, editor. Lettuce, tomato, melon, cucumber, pea,
                                                                                                                  Claim 2: Page 70: 83pp. Eng. 3h
                                                                                                                                                                                                                                                                     Rance I, Gruber V.
                                                                                                                                                                                                                                                                                                             (MERT ) MERISIEM THERATE THE
                                                                                                                                                                                                                                                                                                                                                   22 MAR 1999;
                                                                                                                                                                                                                                                                                                                                                                                         20 MAR 2000; 2000Wo [Brost]?
                                                                                                                                                                                                                                                                                                                                                                                                                             28 SEP 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MO500028808 VI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pisum sativum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plastocyanin dene promoter:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleotide sequence of the promoter MPr1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1007 HSH 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA96,464 standart; INA REFER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 296 BP; 94 A; 74 C 45 C, 83 L, 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rape, beetroot, or sunflower, and monocotyle-bonous species, e.g. wheat, barley, oat, rice, or corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  promoter of the pea plastoc min dene, or comprising a G box operably or functionally linked destroum of a CAAT box. TATA box and transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              petE promoter: chimeric promoter; transgenic plant; MPf1153;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      823 ceasaratetgagesarasaaaa eegatesaratettategeseattetataaaaaa 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               264 adacticall 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     942 Itaattaat 950
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    144 emandatettagenananaanennatenantettateat mattetatadadaa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            84 caetetgtqqcacatetacattatetaaatetaaqecacqteqqaqqataacatattett 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -
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Pred: No. 2.5e 07: 
Pred: Mismatches 25: Indels 29:
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       describes a chimeric expression promoter comprising a promoter of the pea plastocyanic gene, or comprising a data operably or functionally linked upstroam of a CAAU box. TALA box and transcription initiation site. The chimeric promoters are used in expression vectors to producing transcence plants, such as dicetyledonous species, e.g. potato, tobacco.
promoter is delived trom the petE promoter from pea plastracyanib geneby fusing the petE as like and mes enhancer like elements to the promoter MPr1898 (comprising TAIA and CAAT boxes of petE), and then
                                                                                                                                                                                Claim 2: Page 72: 83pp: English.
                                                                                                                                                                                                                                           Chimeric expression promoter for producing diestyledecess and memorialyledonous transferic plants comprises a necessive and sequence derived from a promoter of the pea plastocyanin tene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W02-)0056906 A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleotide sequence of the promoter MPr1143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transgenic plants, such as dicetyledomous species, e.g. potato, tobacce cotton, lettuce, tomato, melon, cucumber, pea, rape, bestroot, or sunflower, and monocotyledomous species, e.g. wheat, barley, out, rice
                                                                                                          The present sequence represents the chimeric presents the philiar. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rance I. Gruber V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MERL ) MERISIEM THERAPECTIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 MAR 2000: 2000Wo IB00+17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pisum sativum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plastocyanin deno promoter; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               petE promoter; chimeric promoter; transgenic plant; Mrc1143;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08 FEB 2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA96466 standard; DNA; 220 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             271 addictedtt 279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       #82 leacaettigidagtetaeaettigalteeettienaacaeataesaaaaaaaaaaaaaaaaaa 94;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        823 ovar aratotspagovaraotaapaa voratova ostotitatosovaritotataapaa eki
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75.2%;
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Bost Local S
Matches 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inserting a G box and fusing a fragment comprising a duplication of the element as2 and as1. A 72 bp fragment comprising the elements as-2, as-2 and as-1 was then deleted. The pett promoter directs cell-specific but not full light-regulated expression in transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operably or functionally linked upstream of a CAAT box, TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous spec.es, e.g. potato, tobacco, colton, lettuce, tomato, melon, cucumber, pea,
                         Claim 9; Figure 6; 45pp; English.
                                                     Enhancer from pea plastocyanin promoter region - used to produce transgenic plants having increased expression of a desired gene
                                                                                                                                     Gray JC,
                                                                                                                                                                                                29-NOV-1995;
                                                                                                                                                                                                                                                                                         WO9720056-A2
                                                                                                                                                                                                                                                                                                                                    promoter
                                                                                                                                                                                                                                                                                                                                                                                 Pisum sativum
                                                                                                                                                                                                                                                                                                                                                                                                                              plastocyanin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pisum sativum plastocyanin promoter region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT77099 standard; DNA; 266
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                                                                                                      WPI; 1997-310610/28
                                                                                                                                                                  (ADTE-) ADVANCED TECHNOLOGIES CAMBRIDGE
                                                                                                                                                                                                                             26-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                            transgenic plant; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-JAN-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT77099;
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                                                                                                                                  Sandhu JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           promoter; enhancer; reporter gene; pea
                                                                                                                                                                                                95GB-0024350
                                                                                                                                                                                                                             96WO-GB02910
                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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81.0%;
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                                                                                                                                   Webster (I;
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Pred. No. 3.2e-07;
0; Mismatches 26;
                                                                      used to produce
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This sequence represents the pea plastocyanin promoter enhances

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                                                                                                                                                                                       17-JUN-1992;
29-JUN-1992;
01-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A.tumcfactors. Expression of the gene promoter of one of more genes of a plant incluses in green or non-green tissues, especially roots, tubers, seeds, stems, flowers or leaves. This enhancer may operate both in normal and reverse directions and can be attached either to the promoter or terminator of the gene to be expressed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence. This enhancer has been used to increase the expression of the GUS reporter gene in tobacco where the GUS gene was under the control of the Pett promoter and 35S CaMV promoter. The enhancer was also used joined to the -330 to +1 region of the paratin promoter PS20 which was joined to the GUS reporter and transformed into potato using which was joined to the GUS reporter and transformed into potato using
                                                                           P-PSDB; AAR44529
                                                                                                                           Xue GP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 266 BF; 102 A; 47 C; 34 G; 83 T, 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                corresponding to nucleotides -444 to -179 of the full length
             food and pulp and paper industries and feed efficiency
                                           New recombinant fungal xylanase – used
                                                                                         WP1; 1994-007529/01
                                                                                                                                                                                                                                                       17-JUN-1993;
                                                                                                                                                                                                                                                                                       23-DEC-1993
                                                                                                                                                                                                                                                                                                                    W09325671-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                Neocallimastix patriciarum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pNPX30 xylanase cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ53480 standard; cDNA; 2503
                                                                                                                                                         (CSIR ) COMMONWEALTH SCI & IND RES ORG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Xylanase, ruminaht animals, fungus, paper, pulp; bagasse:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JUN-1994 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                feedstock; rumen; plant fibre; ss.
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92AU 0003238.
93AU-0003238.
                                                                                                                                                                                                                                                       93WO-AI100294
                                                                                                                                                                                                                                                                                                                                                  /product- Xylanase
                                                                                                                                                                                                                                                                                                                                                                   /*tag-
                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                      . 1935
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67.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 79.2; DB 10, Pred. No. 1.4e=05; M3 smatches 53;
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                          for hydrolysis of xylan for improving ruminant
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Hest Local Similarity 45 7%;
Matches 244; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The cloned xylanase coding sequence is derived from an anaeroble numen fungus. The xylanase has high specific activity for the hydrolysis of xylana. It can be used for fredting pulps in the pulp and paper industry, for freeling handses for more efficient disposal or for the freatment of feedstock to improve nutritional value, denetically modified xylanase genes can also be used for the modification of rumen bacteria to improve plant fibre utilisation by
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2427 adulttaadadtalaalliilitaa adadtaaaqadlalaad=2468
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                   50\% taatqaattqaatqaaqaqttqaattaaaqttqtattaqtaa546
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Search completed: September 10 . 02. 01:04:17 Job time: 8069 see

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result		Query				
No.	Score	Match	Length	ÐΒ	ID	Description
c 1	130 2	13 4	376	=	HF635578	GED 455 28 NFT GEROOF
IJ	115 2	11 9	400	10	HC583521	HCE8=521 BST485233
3	115.2	11 9	480	<u>ن</u>	HC283263	HG581563 ES[4853]5
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œ	101 2	10 4	456	Ģ.	A1.383112	AL383112 MtBC11H02
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AL064082	A1.0668U]		B0582789	AG031915	ALU75432		AL103902	AL108152	AL206908	BEZ49400	ALU61936 Drosophil	F1 240614			AL285149	AL106628		9762361V	FW358135	7:564091	AL069440	BG457563	AL069526	M.632120	A1.06.3921	7. 9.6.5.2.2V	B1271179
Drosophil	Drosophil	Drosophil	EST484535	Pan troql	brosophil	Liddosoud	l indosenti	brosophil	Tet raodon	NF014F06L	Drosophil	30M24 - Sp.6	Drosophil	EST529827	Tet raodon	Drusophil	15.50 th 15.00	brosophi i	CAEa000	Drusophil	Drosophil	NF105G07P	brosophil	NF047A021	brosophil	7221116b	NF050G10F

## ALIGNMENTS

RESULT 1

Source	FEATURES								COMMENT	JOHRNAI	377.11.1		AUTHORS	REFERENCE				ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	HF635678/c
/**cration "Melicaro trupcatula" /*cration "Melicaro trupcatula" /dL.xrel="taxon:3880" /clone="NF109F09DT" /clone_ib="brought" /tissuc_type="Plantlets" /dev_stage "Fooled timepoints" /lote "Vector, Lambda Zapo Contains a mixture of entire plantlets barvested in a series of days post-watering	iceation/Qualifiers	Plate: 109 row: F column: 09 Sey primer: TCACACAGGAACAGCTATGAC.	9	Email: qdmay@noble.orq	Fax: 580 221 7380	2510 San Noble Darkway, Ardmere, OK 73402, USA	The Samuel Roberts Noble Foundation	Plant Biology Division	Contact: May GD	Tippyblished (2000)	Expressed Sequence Tags from the Samuel Roberts Noble Foundation	Flores, H.R., Inman, J.T., Weller, J.W. and May, G.D.	Torrer-Terez, I., Scott, A.D., Harris, A.B., Gonzales, R.A., Bell, C.J.,	l (bases 1 to 376)	Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae;	Spermatophyta; Magnoliophyta; endicatyledons; core cudicots:	Eukaryota; Viridiplantae, Streptophyta, Embryoph,ta, Tracheophyta:	Medicago truncatula	barrel medic.	EST.	BF635678.1 G1:11899836	HF635678	5', mRNA sequence.	DT1F1077 Drought Medicago truncatula cDNA clone NF109F	BF635678 376 bp mRNA linear EST 19-DEC 2000	

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17 K.18 42.71
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Bukaryota; Viridiplantac: Streptophyta: Embryophyta: Dacheophyta:
Spermatophyta: Mauno.sophyta; cudicotyledons: core endicots;
Rosidae: eurosids 1: Pabales; Fabaceae; Publicacideae; Tritoliege
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Seg primer: SKmed (CSA 4AA PTA 4ff 4AL PP).
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University of Minnes and
411 Horland Hall, 1901 Upper Butord Circle, St.Paul, MN 55108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (2001)
Contact: Carroll P. Vince
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of D., Van Akon S., Ottoberek,E., Ch. J. and Frason,C.M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: vance004 maross.te.unn.edu
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                                                                              Zhoic "Vector: pBluescript SK; Site_1: Brokh; Site_2: Xhoi: cPNA was prepared from polyA+ enriched RNA from ellectrice root modules harvested one month post
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post inwentation with SmorthFobium melitoti"
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inoculation with Sinochizobium meliloti. The cDNA was literthonally ligated into the Uni ZAP XP yester from Stratagene and packaged using digapack life Orld packaging
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University of Minnesota name: M48464le [15]R sequence name
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Tel: 612-625-5715
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48 c. 73 g. 153 t.
                                                                              post inoculation with Sinorhizobium metalori"
Zlab host "E. coli strain XIzeke"
Znote-"Vectur pBluescript SK: Site I: Ecoki
Xhol: CDNA was prepared from polyA+ enriched RNA from effective roof modnics buryested one month post inoculation with Sinorhizobium metileti. The cONA was
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/ALXEST "GEOGRAPH"
/Alone "Java /APIL"
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                                                                                                                                                                                                                                                                                                             University of Minnesota name: M261199e
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Tel: 612 625 5715
                                                                                                                                                                                                                                                                                                                                                                                                                         Department of Agronomy and Plant Genetics University of Minnesota
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Carroll P. Vance
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26 0 75 g 179 r
                                                                                  /organism="Medicago truncatula"
/Cull.var="qenotype A17"
/db_xre1-"haxon:3880"
/clone="pGVN-58A15"
                /tissuc_type="N2 fixing root nodules"
/dev_stage="effective root nodules harvested one month
                                                               /clone_lib="GVN"
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EST484001 GVN Medicago truncatula conA clong peva 69021 50 end,
                                                                                                                                                                                                                                                                         Tel: 612 625 5715
Fax: 651-649-5058
                                                                                                                                                                                                                                                                                                                                                          Department of Agronomy and Plant Genetics University of Minnesota
                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2001)
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                                                                                                                                                              MTCDS95TK More information is available at: http://www.medicago.org Seq primer: SKmod (CTA gAA CTA gtg gAT CC).
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                                                                                                                                                                                                                                                                                                                             411 Borland Hall, 1991 Upper Buford Circle, St.Paul, MN 55108 USA
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Spermatophyta: Magnoliophyta: eadleotyledoms: core endicots:
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/note-"Vector: pBluescript SK:; Site_1: EcoRI; Site_2:
Xhol; cDNA was prepared from polyA: enriched RNA from
effective root nodules harvested one month post
/cultivar-"genotype A
/db_xref-"taxon:3880"
/clone="pGVN-69021"
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                                                      /organism-"Medicago truncatula'
/cultivar="genotype A17"
                                                                                                                                       Location/Qualifiers
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/tissue\_type="N2-fixing root modules"

/clone\_lib-"GVN"

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Plant Physiol. (17 (4): 1825-1882 (1998)
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Covity.P A , Smoth.P 3. and Long,S.R
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Spermatophyta: Mauneliophyta: esdiestylekas: sore esdiests)
Rosidae: eurosids I: Fabales: Fabaceae: Papilionoideae: Tritolieae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Department of Hiological Sciences and Howard Hughes Medical
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                                                                                                                                                                                                                                                                              Email: ta.srl storsyth; _stantord.edu
                                                                                                                                                                                                                                                                                                                     Pax: 650 725 8309
                                                                                                                                                                                                                                                                                                                                                Tel: 650 724 (2)
                                                                                                                                                                                                                                                                                                                                                                       Gilbert Biology, Stantord, CA 94/05 Sazd, USA
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Citions "pp. "East laris & "Tp."
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elloctive root modules harested one menth post
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post inoclation with Sinorhizobium mediloti"
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Best Local Similarity (4.4%) Pred. No. 1, center
Matches [164] Conservative (0) Mismatches (6); Indels
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                                                                                                                     betermination of this BAY end sequence was carried out as part of a collaboration with the BAY end sequence was carried out of part (BDAY). The BDAY is construct to a physical morphical denome Project (BDAY) in melanosaster denome using these BAYS. For further information picture see http://www.fruitfy.org.The BAWE frosophila melanosaster BAY (ibbary was prepared by Kirottoyo oscordawa and Aaron Mammoser in Pleter de lond's Laboratory in the begarinent of Concer denotics at the Koswell Park cancer institute in Bultato, NY. The library is named BAY 98 and was constructed by partial EcoRI direction of brosophila DAA provided by the BADE isoderne strain y2, or bw sp., the same strain sed for the BADE's Pl and EST libraries. A more detailed description of the Ibrary of the control of the BADE is the construction transfer and bow to order individual BAY clones. The control Unitary, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (02-JHN 1999) Genoscope Centre National de Sequencade HP 191 91006 EVRY cedex FRANCE (E-mail: segret genoscope.cns.tr - Web : www.genoscope.cns.tr)
                                                            filters for hybridization from the BACPAC Resource Center can be found at http://www.pac.myd.huttalo.cd./dosephila bacching
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Pterydota; Neoptera; Endopterydota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosochila
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AL069706
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from edougating root bairs (50% w/w) and 2 com not typs
(70% w/w). Xhol olique di linker primer and ECOK:
adapters were ased, cBMAs was cleared unidirectionally
into lambda ZAP Express (Stratanene), amplified, and
mass exclased into pHK CMV vector plasmids. More
information is available of the p.//bio.sRLM.stant.cd.epu.
a 156 c 88 q 179 i he p.//bio.sRLM.stant.cd.epu.
towation/Qualitiers
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                     Journet,E.P., Crespeau.H. van-Tuinen.D. Goury T. Jaillon.o.
Niebel,A., Carreau,V., Chatagnier,O., Kahn,D., Gianinazzi-Pearson,V. and Gamas,P.
                                                                                                               Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I; Fabales, Fabaceae, Papilionoidman: Trifolioae;
                                                                                                                                                       Medicago truncatula
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
                                                                                                                                                                                                                                           AL383112.1
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Medicago truncatula ESTs from endemycorrhizal roots
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/db_xref="taxon:7227"
/clone_lib="RPCI-98"
/clone="BACR29B23"
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38.0%; Fred No 2e.06
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Biologie Moleculaire des Relations Plantes-Microorganismes,
CNRS-INRA, BP 27 31326 Castanet-Tolosan Cedex, France (Email .
Medicago truncatula
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BP 191 91006 EVRY cedex - France
Email: segref@qenoscope.cns.fr, Web : www.genoscope.cns.fr
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                             barrel medic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             extracted from whole root systems, cDNA was prepared from FolyA: enriched RNA. The cDNA was directionally ligated into Uni-rap XB vector from Stratagene and packaged using Gigapack Gold packaging extracts. Plasmids containing cDNA claserts were mass excised from phage stocks. Ging ExAssit helper phage and propagated in SUR cells. Close ordering and sequencing was performed by the Centre National de Sequencage (Cenoscope, Evry, France). Note: ESI may be of tungal origin."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              72b at 25 C, before transplanting into a 1/3 Epoisses soil: 2/3 calcined Terragreen mix in the presence of onion root fragments colonized by the arbuscular mycorrhizal
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Journett E Po Stessent Hoo van Tuinen. Do Gerzy, Joj Jaillon, Oo,
Niebel, A., Garreau, V., Chatagnier, co., Kahm, Do, Glaminazzi Pearson
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t : Pascal Gamas and Effenne Pascal Journet, Laboratoire de
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Texas AM University

Department of Plant Pathology and Microbiology, Rm L20 L1F

Consequence of the Collogo Station. IX 7/84% 2107, USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              315 aaatagttgtacaaatatcatt 336
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                                                                                                                                                                                                                                                                                                                                                                     AL368181 MtBA Medicajo truncatula cDNA clone MtBA22F11 T7, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Department of Plant Pathology and Microbiology, Rm Peterson Bidg, College Station, TX 77843-2132, USA Tel: 409 845 8743
Fax: 409 862 4790
Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
                                       Contact: Genoscope
                                                     Medicago truncatala ESTs from nitrogen starved roots unpublished (2003)
                                                                                                             Journet, E.P., Craspeau, H., van-Tuinen, D., Gouzy, J., Jaillon, O., Niebel, A., Carreau, V., Chatagnier, O., Kahn, D., Gianinazz, Pearson
                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta, Magnoliophyta, Audiontyledons; Pore endionts; Rosidae; Eurosids I, Fabares; Fabaceae, Papillonoideae; Trifolieae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: dcook@ppserver.tamu.edu
Seg primer: SQ-HAG-S (AMAGAGGAAAGAGGTATGAGGATGATTAGG)
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Contact: Cook DR
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Medicago, truncatula BACs linked to the DOMI (DMII) locus
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                                                                                                and Gamas, P.
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/db_xref-"taxon:3880"
/clone="183D}7"
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Locus
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                                                                                                                                                                                                                                                                                                                                                                                                                                     T132103b Medicago truncatula BAC library Medicago truncatula genomic clone 10F20, DNA sequence.
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Contact: Cook DR
                                                                                                                                                                                                              Eukaryota: Viridiplantae; Streptophyta; Embryophyta; Iracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
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Contact : Pascal Gamas and Ettenne-Pascal Journet, Laboratoire de
Hiologie Moleculaire des Relations Plantes-Microorganismes,
CNPS: (NPA, NP 27 31326 Castaset Telesa: Cedex, France (Email :
Texas A&M University
                       The Crop Biotechnology Center
                                                                                                     BAC survey sequencing of Medicago truncatula
                                                                                                                                  1 (bases 1 to 492)
Kim,D., Peng,H., Ellis,L.
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//dev_stage="barvested after 3 days of Nestarvation"
//note="Weetor: pBlueseript pSK; Site_1: FeoR; Site_2:
Xhol; Plants were grown in an aeropoine chamber for 14
days on nitrogen=rich medium followed by 3 days on N free
medium. RNA was extracted from root tips (1-3 cm), CDNA
was prepared from polyA+ enriched RNA. The cDNA was
directionally ligated into Unicapak vector from
Stratagene and packaged using diapak vector from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and propagated in SMLR cells, Clone ordering and sequencing was periormed by the Centre National de Sequencage (Genoscope, Every, France)."

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                                                                                                                                                                                                    JOHNA!
                                                                                                                                                                                                                                                                      AUTHORS
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T137727b shotom sub-library of BAC close 1988 Medicago truncatula
genomic clone 10 F 20 071, DNA sequence.
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other name: DKO4 kk; date: BZ20/9B; Shbmilled to the Dalabase of
othernome Survey Sequences (GSS) on 09.7% or More into matice is
available at 'http://chrysre.tamu.edu/medreado';
Seq primer: SQ FAC k (ACA/ACGAAA/AGCTAFGACCATGATTACG)
Class: BAC ends.
                            Texas AAM University
Department of Flant Esthology and Microbiology, Rm 120 L.F.
Deterson Bldg. College Station, TX 2744 C21 C2 USA
                                                                                                                                                                    Contact: Took DR
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                                                                                                                          The Crop Biotechnology Center
                                                                                                                                                                                                                            BAC survey sequencing of Medicago truncatula (1999c)
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(thet name: BSC 20-71; data; 8/29/99; Submatted to the laterase General Sarvey Sequences (CSS) as my/20/99; More information is also in the post of the control of the Sequences (CSS) as a fight to the control of the Sequences (CSS) as a fight of the control of the Sequences (CSS) as a fight of the control of the Sequences (CSS) as a fight of the control of the Sequences (CSS) and the control of the control 
cancer denotics at the Roswell Park Parcer institute in Buttare, NY. The library is named RPCI-98 and was constructed by partial Record disestion of Drossphila DNA provided by the BOGP from the Alacetic strain 25. In the Sp. The dame strain used for the HBMP's PL and EST libraries. A more detailed description of the library, or dual how to order individual HAC clones, the entire library, or
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                                     513 tgatgaaagagttggattaaagttgtatt 541
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                                                                                                                                                                                                                                                                               333 cattgaggaatttgacaasagctacacaaataagggttaattgctgtsaattaaatsaagga 392
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     843 ADDRWDAWAWKWDDAWAWGAPTADPROWGDRAGKPGGARKPPDPKRADDKRDAADDRDDA 784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        903 ADDDTDGTKUDDBKKKUDJWDKAKGTWGDATWAWAATDWWWWGWADADWWTWDAAADDWW 844
                                                                                                                                                               603 TWIDARKADROWAKARAWRARPORARAARADRRWIIKGKIITATWIIWAARAAWWAWWAW 544
                                                                                                                                                                                                                                                                                                                                                                                                                   723 ККІЛЬМІЛАПАЛЛІАКІЛЬКЬУ КУКЛІЛІЛАПАЛКОККІ СРКЬБЬКОКАТИ ІКПЛАММАЛЛАМИ 664
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Location/Qualifiers
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APPLICANT: Miller, Louis H. APPLICANT: Peterson, David S. APPLICANT: Wellems, Thomas E. APPLICANT: Wellems, Thomas E. TITLE OF INVENTION: BINDING DOM TITLE OF INVENTION: BINDING DOM TITLE OF INVENTION: AND PLASMOT NUMBER OF SEQUENCES: 45 COMPRESSEE: KNOBBE MATTENS OLS SIRBEL: 620 NEWPORT CENTER ADDRESSEE: KNOBBE MATTENS OLS SIRBEL: 621 NEWPORT HEACH STATE: California COUNTRY: US ZIP: 92660 COMPUTER: IBM PC COMPATIBLE COMPUTER: IBM PC COMPATIBLE COMPUTER: FORM: MEDIUM TYPE: Ploppy disk COMPUTER: FORM: PC-DOS/MS- COMPUTER: BLEATION DATA: APPLICATION NUMBER: US/08/487 FILING DATE: 10-SEP-1993 CLASSIFICATION NUMBER: US/08/487 FILING DATE: 10-SEP-1993 ATTORREY/AGENT INFORMATION: NAME: ISFAELSEN, NEE  REFERENCE/JOCKET NUMBER: U-655 REFERENCE/JOCKET NUMBER: U-656 REFERENCE/JOCKET NUMBER: U-657 INFORMATION FOR SEU ID NO: 13: SEGUENCE CHARACTER:STICS: IFENTAL 19124 LASC PAIRS TYPE: NOTALESTICS: IFENTAL 19124 LASC PAIRS MYSCHIELETAN: 19124 LASC PAIRS MYSCHIELETAN: US/08/487 MYSCHIELETA	28 45 4.6 1186 2 US-08- 24 44.5 4.5 251 4 US-08- 25 44.4 4 5 3529 4 US-08- 31 44.4 4 6 3529 4 US-08- 32 44.4 4.6 4383 6 517730 34 44.4 4.5 9636 1 US-08- 35 44.4 4.5 9636 1 US-08- 36 44 4.5 9636 1 US-08- 37 43.6 4.5 9636 1 US-08- 38 43.4 4.5 9636 1 US-08- 39 43.4 4.5 9636 1 US-08- 44 4.5 9636 1 US-08- 45 43.4 4.5 9636 1 US-08- 46 43.4 4.5 9636 1 US-08- 47 42.8 4.4 483 6 51750 41 42.8 4.4 483 6 51750 42.8 4.4 4983 6 51750 42.8 4.4 4983 6 51750 42.8 4.4 4983 6 51750 42.8 4.4 8920 2 US-08- 45 42.8 4.4 8920 2 US-08- 45 42.8 4.4 8920 2 US-08- 45 42.8 4.4 8920 4 US-09- 46 42.8 4.4 8920 4 US-09- 47 42.8 4.4 8920 4 US-09- 48 42.8 4.4 8920 4 US-09- 49 42.8 4.4 8920 6 E23116 48 4.7 893827 49 41
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                                                                                                                                                          Sequence 14, Application PS/08.332463 Patent No. 5670367
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APPLICANT: SCHEFFLINGER, F. APPLICANT: FALKNER, F. A. TITLE OF INVENTION: RECOMBI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                284 tgagagagaaagtigtacaaaaittgtaccaaaatagttgtacaaaatatcattgaggaat 343
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   44 aaaagtttaagttagcaagtgt-placatttttatttgaacaaaaataftcacctactact 103
                                                                                              DORNER, T
   RECOMBINANT FOWLPOX VIRUS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
6.8%; Score 66.4; DR 1: Length 2218;
Best Local Similarity 9.8%; Pred. No. 1.4e-05;
Matches 46, Concervative 229, Minualshee 195, Indele 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 14:
IMMEDIATE SOURCE:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
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TERMITHS SYSTEM. FC 10-8,785 DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
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                                    372 atturtutaaataaataaggatgarurattagagagatgtarrattagagaattttiger 431
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                                                                                                                  252 gaaagaggaadaataaaaaassataatytaagtutgaguaaguuugttgtacaaaagttgta. Kil
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CTTY: Alexandria
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/232,463
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLONE:
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US-(8-998-416-288
                                                                                                                                                                                                                                                Query Match 6.8%; Score 65.8, DB 4, Length 837; Best Local Similarity 46.7%; Pred No. 1 4e-05; Matches 279; Conservative 0; Mismatches 312, Indels
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APPLICANT: Philip
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APPLICANT:
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APPLICANT: Knechtle, Prilipp
APPLICANT: Rebischung, Corinne
TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHBYA COSSYPII
TITLE OF INVENTION: AND USES THEREOF
NUMBER OF SEQUENCES: 1152
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
URIGINAL SOURCE:
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| 180 tttgttgcaacatttgayaaaattttgttgttctctcttttcattggtcaaaaacaata 239
                                                                                                                       120 aaacattagagtaaagaaatatggatgataagaacaagagtagtgatattttgacaacaa 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1 0, Version #1 40
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CITY: Research Triancle Park
STATE: No. 6239264th Carolina
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                                                                                 66 ATTAAATTATTTAAACTATTATTATCATTATTTAATAAATTAATTATLIGATTATTAATA 125
                                                                                                                                                                                                        60 aagtgtgtacatttttat::tgaacaaaaatattcacctactgttataaatca:tatt 119
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                                                                                                                                                                   6 AAATATATATAATTAATTAATTAAAGATTAATATAAAACTTTTTTATIATAATATTTAAGT 65
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nucleic acid
NDEDNESS: single
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Steiner, Sakine
Mohr, Christine
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                  INFORMATION FOR SEQ ID NO:
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APPLICANT: Pohlmann, Rainer
APPLICANT: Steiner, Sabine
                                                                   TELECOMMUNICATION INFORMATION:
SEQUENCE CHARACTERISTICS:
                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    246 AAAATATTATTAGATATTATTTTTTTTAATAAAITATTAAATAGATTATCAATAATT
                                   TELFFAX:
                                                                                    REGISTRATION NUMBER: 38, REFERENCE/THANKET NUMBER:
                                                                                                                                                                                                                           FILING DATE: 24-DE
                                                                                                                                                                                                                                                                                   SOFTWARE:
                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                     COMPUTER:
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CITY: Research Triangle Park
                                                  TELEPHONE:
                                                                                                                           NAME:
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                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE:
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                                                                                                                        Meigs, J. Timothy
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                                   3: 919-541-8587
919-541-8689
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VENTION: GENOMIC DNA SEQUENCES OF ASHBYA GOSSYPII
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Knechtle, Philipp
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Patent No. 62/9264
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                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                 APPLICANT. RODISCHURG, COLIGION SEQUENCES OF ASHRYA COSSYPIL TITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                APPLICANT: Poblimand, Rather APPLICANT: Steiner, Sabine APPLICANT: Mohr, Christine APPLICANT: Wendland, Jursen APPLICANI: Knechtie, Philipp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORIGINAL SOURCE
                      CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6239264artis Corporation
                                                                                  NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEC 10 NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 4) DEC 1996
ATTORNEY/AGENT INFORMATION:
NAME: MOIGS, J. Timothy
REGISTRATION NUMBER: 38, 241
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OKICINAL SOURCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
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488 tradaccatqtqaffatftadfqadftqaffqadftqqaffaaqffqfatfaqfaa 546
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3) agreementaget grant of a calabate of the agreement for a calabate of the ca
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19) cutttgagaaaattttgttgttctctcttttcaftggtcaaaaacaatagagagagaaaa 280
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COMPUTER: THM PC compatible
OPERATING SYSTEM: PC:DOS/MS Des
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STATE: No. 6239264th Carolina
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EDNESS: single
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                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 636 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 919-541-8587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1 0, Version #1 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTURNEY/AGENT INFURMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: GEROMIC DNA SEQUETITIE OF INVENTION: AND USES THEREOF
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                                                 180 ttttgttgcaacatttgayaaaattttgttgttcttctttttcattgqtcaaaaacaata 239
120 aaacattagagtaaayaaatatggatgataagaacaagagtagtgatattttgacaacaa 179
                                                                                                                                                                                                                                                    y Match 6.1%; score 58.8; DH 4; Length 636; Local Similarity 45.5%; Pred. No. 0.00036;
                                                                                                                                                                                    TELEPHONE: 919-541-8689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME. Meigs, J. Timothy
REGISTRATION NUMBER: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: CH 0016/97 FILING DATE: 31-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                 TOPOLOGY:
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                                                                                                                                                                 6 AAATATATATAATTTAAT UTATAAAGATTAATATAAACTTTTTTTATTATAATATTTAAGT 65
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3054 Cornwal is Road
                                                                                                                                                                                                                                    Conservative
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VENTION: GENOMIC DNA SEQUENCES OF ASHBYA GOSSYPII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wendland, Jurgen
Knechtle, Philipp
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Steiner, Sabine
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N: 435
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Query Match
Best Local Similarity
Matches 226; Conserv
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Sequence 1, App...
Sequence 1, App...
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                                                                                                                                                                                                                                                        TELEFAX: 703-816-4100 INFORMATION FOR SEQ ID NO:
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APPLICANT: Flores, Maria V
APPLICANT: O'SHilivan, William J
                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
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TITLE OF INVENTION: phosphate synthetase II
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  246 AAAATATTATTAGATATTATTATTTTCTTTAATAAATTATTAAAIAGATTATCAATAAIT 305
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                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                    STRANDEDNESS: single
                                                                                                                                                                                              TYPE: nucleic acid
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                                                                                                                                                                                                                LENGTH:
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                                                                                                                                                                                                                8920 base pairs
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1100 No 584957∛th Glebe Poad, 8th Floor
      Conservative
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  Score 57.2; IB 2; Length 8920;
Pred. No. 0.0012;
0; Mismatches 263; Indels 1
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                                                                                                                                                                                                         Ouery Match 5.0%: Score 57.2; DE 4; Longth 8920; Hest Local Similarity 46.1%: Fred. No. 0.0012;
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09 | 150 | 741 | 1
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NUMBER OF SEQ ID NoS: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Stewart et al
                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13, Application US/08487826H Patent No. 5993827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION
                                   INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENCTH: 19124 base pairs
                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sim, Kim L.
APPLICANT: Chitnis, Chetan
APPLICANT: Miller, Louis H
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: BINDING DOMAINS FROM PLASMODIUM VIVAX
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                                                                                                                                                                               NAME: ISTACLSON, Ned
REFERENCE/DOCKET NUMBER: 29
                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                             TELEFAX:
                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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  STRANDEDNESS:
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(619) 235-0176
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                                                                                                                                                                                                                           US-08-998-416-1137/c
                                                                                                                                                                                     Sequence 1137, Application US/08998416 Partent No. 6239264
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                                                                                                                                                                 GENERAL INFORMATION:
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      TITLE OF
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T: Rebischung, Corinne
INVENTION: GENOMIC DNA SEQUENCES OF ASHBYA GOSSYPII
                                                                               Pohlmann, Rainer
Steiner, Sabine
Mohr, Christine
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                                      Wendland, Jurgen
Knechtle, Philipp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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COMPUTER: HOPPY disk
COMPUTER: POPPY
OPERATING SYSTEM POPPYS/MS-POS
SUFTWARE: PatentIn Melease #1.0, Version #1.
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 636 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/TOO'KET NUMBER PR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                              483 TAAAGTTAAAAFTAAFTTTAAFAAFAAFAATATAAAAAGAFFAAAFAAFAAFAAFAAC 424
441 aaaagaaagaataaattattittaaaattaaaaqttgaytcattigaitaaacatgigat 500
                                           243 TAACTAATTTAAAATTTGAACATAGACTAAATAGTATTCATATTAAATATTATTTTATA 184
                                                                                                                                                                                                                                                         264 ataaaaacataatgigagtatgagagagaagttglacaaaagttgtaccaaaatagttg 929
                                                                                                                                                                                                                                                                                                                                                                                            423 ATAATATTATAAAATAGATATTATAATAAAAAAAATAATTAATTTACAATATTTAAATAAAT\rightarrow64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     324 tac---aaatateattgaggaatttgacaaaagetacacaaataagggttaattgetgta 380
                                                                                                                                                                                                                    303 TTATTGATAATCTATTAATAATATTTATTAAAGAAAATAATAATATATCTAATATTTTAA 244
                                                                                                                                                                                                                                                                                                       144 atgataagaacaagagtagtgatattttgacaacaattttgttgcaacattttgagaaaat 203
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REGISTRATION NUMBER:
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STRANDEDNESS: single
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              Query Match
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 Best Leval Similarity
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Patent No. 5476/8]
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APPLICANT: Hall Richard L
APPLICANT: Grund! Month
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ATTORNEY/AGENT INFORMATION:
                                                                                                          PEATURE:
                                                                                                                                                      LOCATION:
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                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 1511 base pairs
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APPLICATION NUMBER:
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COMPUTER: IBM PO compotition

ODERATING SYSTEM: PO LOCAL MEDIUM

SUPTWARE: Patentin Release #1.0, Version #1 25
CURRENT ApplifONTION DATA
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Score 52: [58 ]
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APPLICANI: Hall, Richard I.
APPLICANI: Grand: Michael E.
TITLE OF INVENTION: No. 57218526] Entomagostic. Expression System
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                                                                                                                     COMPUTER READABLE FORM
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MEDIUM TYPE: Floppy disk
COMPUTEE: IBM PC computible
OPERATING SYSTEM: PC-1005/MS DOS
SPETWARE: Futentin Release #1.0, Version #1
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: 19-AU

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ATTORNEY/AGENT INFORMATION:
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ORIGINAL SOURCE:
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1035 AAAAGTTTATATTTTCAAATTTAAATATTATTAATTTAAATTTTATAACAAAATTAAAA 1094
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Local Similarity 44 7%;
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                                                                    GAAGAATTGCATATAATATATTATGATAATAATATTTTAAATAATATTCCAGAAAAATATT 1034
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EDNESS: double
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0; Mismatches 355, Indels 13, Gaps
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Patent No. 5935777
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                                                                                                                          NEORMATION FOR SEQ ID NO:
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APPLICANT: Hall, Richard L.
APPLICANT. Guidl, Michael F.
TITLE OF INVENTION: No. 5935777+1 Entomopoxvirus Expression System
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APPLICATION NUMBER: US 0:
FILING DATE: 19-FEB-1941
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                                                                                                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 904-375-8100
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APPLICATION NUMBER: WO 9:
FILING DATE: 12-FEB-1992
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  MOLECULE TYPE:
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OPERATING SYSTEM: PC-DOS/MS-TANS

SOPTEMARE: Palentin Boloci
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                     TOPOLOGY:
                                     STRANDEDNESS:
                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER -
                                                                                                                                                                                                                               NAME: Bencen, Gerard H
REGISTRATION NUMBER: 3
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DEDNESS: double
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2421 N.W. 41st Street, Suite A-1
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DNA (genomic)
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                                                                                                                               RESULT 14
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Matches 298; Conservative
                          GENERAL INFORMATION:
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APPLICANT:
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Philippson, Feter
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Pred. No. 0.011:
d: Mismatches 455; Indels 13; Gaps
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APPLICANT: Knichtle, Philipp
APPLICANT: Rebischund, Colium
TITLE OF INVENTION: GENERAL DAN SEJÜENCES OF ASHBYA (1888YELL
VUMHER OF SEQUENCES: 1152
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402 AHTATAAAAAATAATAATHACAATATHAAAATAATTAAAADTHEATAATAATAATAAAA 343
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                                                                                                                        282 tätdagagadaagittolaraaaaditgiacraaaatagitgiacaaatattoloota eetooligaga (4)
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                                                                                   342 attitgacaáaagetaeaeaaataagggitaattgetgtaaataaataaggatgaegeatt 48]
                                                                                                                                                                                          REGISTRATION NUMBER: 48
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Research Triangle Park
: No. 6239264th Carolina
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Mohr, Christine
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Patent No. 6239264
                                                                                                                       Matches
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                    MOLECULE TYPE:
ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION: NAME: Meigs, J. Timothy
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ADDRESSEE: No. 6239264artis Corporation
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TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHRYA GOSSYPIT
TITLE OF INVENTION: AND USES THEREOF
NUMBER OF SEQUENCES: 1.52
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142 ggatgataagaacaagagtagtgatattttgacaacaattttgttgcaacatttgagaaa 201
                                        462 ttaaaattaaaagttgagtcatttgattaaacatgtgattatttaatgaattgatcaaag 521
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CITY: Pesearch Triangle Park
STATE: No. 6239264th Carclina
                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
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FILING DATE: 31-DEC-1996
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Knechtle, Philipp
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Pohlmann, Rainer
Steiner, Sabine
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235 AAAATAATAAAATAACAAAATAAT 212
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Search completed: September 1, 2002, 00:57:51 Job Lime. 7943 Sec

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Maximum DB seq length: 2000(0000
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Perfect score:
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Copyright (c) 1993 - 2000 Compugen Ltd
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## ALIGNMENTS

REFERENCE ATTHORS TITLE DEFINITION ACCESSION VERSION RESULT Ax105269 LOCUS KEYWORDS SOURCE ORIGIN BASE COUNT FEATURES JOURNAL. ORGANISM source synthetic construct synthetic construct artificial sequence.
1 (bases 1 to 731)
Vezina, LP and D'Apust, M.A. AX105269 731 bp Sequence 3 from Patent Wo0125455. AX105269 Promoter for regulating expression of foreign genes patent: WO 0125455-A 3 12-APR-2001; AX105269 1 GI-13921419 Medicago Inc. 299 a expression" /organism="synthetic construct" /db\_xref="taxon:32630" /mote "Sequence to be used as a Fromoter for requiating Location/Qualifiers (CA) 109 q 210 ( ୍ NA lincar PA" 30 APR 2001

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Vezina, L.F. and D'Acust M A
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                                         l (bases 1 to 834)
Rance, I., TheLsen, M. and Gruber, V.
Patent FP 2741358-A 1 24 SEP 2000;
MERISTEM THERAPEUTICS (FR)
                                                                                                                synthetic construct.
synthetic construct
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Ranco, 1., Thoisen, M. and Grubor, V.
Patent: FR 2791358 A 2 29 SEF 2000;
MERISTEM THERAPEUTICS (FR)
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Rosidae; eurosids i, Fabales; Fabaecae, Papilionoideae; Vici
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/product="plastocyanin"
/proteir_id="CAA48393.1"
/db_xref="GI:4379383"
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/organism-"Pisum sativum"
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                                                                             tyooccustayaytoayttaactoatttittätätittoalayatcaastaayayaaataace, 416
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/product "mature plastocyanin (AA 1-99)"
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/db_xrof-"G1:20846"
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/db_xrof-"SNISS-PROT:P16002"
/translation-"MAIVESTIVATESFSGLKINAATKVSAMAKTPISTSQSPRLCVR
ASLKTPTVALVATAASAVLASAALAVEVIT-ASI***ILAFVPS-15V-ACFTIVEKNNA
GFPHNVVFDEDEIPAGVDASKISMPEEDLLNAPGETYSVKLDAKGTYKFYCSPHQGAG
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                                                                   MERISTEM THERAPETITIS (FR)
                                                                                                       Rance, i.. The idea. M. and Graber, V_* Patent: Fig. 2/91 for A \leftarrow 29 SEP 20001;
                                                                                                                                                                              artificial sequence (bases 1 to 200)
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/db_xrct_"taxon:32630"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match (12.2%) Score 89.4; DB 6; Leggth 281; Best Local Similarity (7.8%; Prod. No. 1.8e-05; Matches (4); Conservative (0); Mismatches (6); Indets (2.28)
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AX036599
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auce, L., Theisen, M.
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promoter MPr1108"
a 71 c 45 g 80 t
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/db_xret-"taxon:32630"
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                                                                                                                      1 (bases 1 to 296)
Rance, I., Theisen, M. and Gruber, V.
Patent: FR 2791378-A 9 29-SEP-2000;
MERISTEM THERAPEUTICS (FR)
                                                                                                                                                                                                synthetic construct
synthetic construct
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                                                                                                                                                                                                                                                       Sequence 9 from Patent FR2791358. AX035603
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1 73 c 42 g
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         /note-"Promoter MPr1111 created by inserting at -99 bp position of MPr1098, an 18 bp element containing a 'G' box and fusing a sequence of 58 bp (duplication of the element
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CaMV promoter
                                                                  /organism="synthetic construct"
/db_xre:="taxon:32630"
                                                                                                             Location/Qualitiers
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76.2%;
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Pred No 2:14-05;
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1 (bases 1 to 303)

Rance, I., Theisen, M. and Gruber, V.
Patent: FR 2791358-A 8 29-SEP-2000;
MERISTEM THERAPEUTICS (FR)
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Sequence 6 from Patent PR2791358
AX0366602.1 SI:11226197
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                                                                                                                                                                                                                                                                                                                              promoter MPr1153"
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76.2%;
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Search completed: September 1, 1802, 00:55:25 Job time: 17711 see
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        DB 6;
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                                                                                                                                                Ranco, L., Theisen, M. and Gruber, V.
Patent: FR 2791358 A 10-29 SEP 2000;
MERISTEM THERAPEUTICS (FR)
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Sequence 10 from Path-of FR2791458.
AX0.66.04 AX0.66.04.1 G1:11.226.29
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promoter MPrIII43"
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Result No.	Score	Query Match Length DB	Length	DB	ID	Description
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Novel promoter for regulating expression of foreign genes organisms, more specifically in a leaf-specific manner in plants :

in transgenic

WP:, 2001-266316/27.

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## ALIGNMENTS

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AASUZIZB standard; DNA; 731 BP

04-0CT-1999; W0200125455-A2 Synthetic. Promoter: transperie plant, no no odyleden, die tyleden, gjøre sperm; ds Fromoter #3 for regulating expression of foreign genes. AAS02128; Vezina L, (MEDI-) MEDICAGO INC 02-00:-2000; 2000wo-ca01144 12-APR-2001 18-JUL-2001 D'Aoust M; (first entry) 99US-0157129

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AAS02127

AASO2127 standard; DNA; 9/1

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Ouery Match 100.0%; Score 731; DB 22; Best Local Similarity 100.0%; Pred. No. 6.6e 133;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The sequence represents the coding sequence of promoter #2 for regulating expression of foreign genes in transagnic organisms. The promoter is useful for regulating the expression of foreign genes in transagnic organisms, particularly plants, e.g. mespectyledoms, directyledoms or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 9; 9pp; English.
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                                                                                                                                                                                             Novel promoter for regulating expression of foreign genes in transgenic organisms, more specifically in a leaf-specific manner in transgenic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             are used in expression vectors for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, tape, lectroot, or Suntlower, and monocotyledonous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression promoter comprising a promoter of the psa plastocyanin gene, or comprising a G box operably or functionally linked upstream of a CAAT box. TATA box and transcription initiation site. The chimeric promoters
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transgenic tobacco plants—the promoter is used to construct chimeric
promoters of the invention, the specification describes a chimeric
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                    peth promoter: chimeric from 'er; transdenic plant
                                                       Nucleotide sequence at the perE promoter from plastocyanin gene
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Nucleotide sequence of the promoter MPT1096

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Best Local Similarity 82.8%; Pred. No. 1.7e 28;
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Best Local Similarity 84.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       barley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MERI-) MERISTEM THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  508 caatootyatyayataacoomittigageeen geachetyiggeaminiamattatet 567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    448 tatatttactaaaaaatclaaqccacgtaqqaqqataacatccaatccaaccaatcacaa 507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39 tatttactaaaaaaatctaagnnanginggaggataanatiicaaccisagecaatsacag 98
                                                                                                                                                                                                                                                                                                                                aattcagtagagaagaaactcatt 302
                                                               aagagaagagactaattaattaat 710
                                                                                                                                      coattictataaaaaatoacottioligtigtictictictit tiogaliticocti caaacacataca
                                                                                                                                                                             aaateacatattetteeaeacatettageeaeacaaaaaeeeaateeseatetttateat 218
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       beetroot, or sunflower, and monocotyledonous species, e.g. wheat,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000WO-1B00317
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 Mismatches

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Pred. No. 4.1e-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            38 C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            86 T, 0 other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 21.
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RESULT AAF 96460

AAA96462

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AAA96462 standard; DNA; 250 HP

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                                                                                                                                                                                                                                            Best Local
                                                                                                                                                                                                                                                                                                                                                                bux, TATA bux and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, beetroot, or sunflower, and monocotyledonous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   minimal reference promoter on the petE promoter. The petE promoter directs cell-specific but not full light-regulated expression in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric expression promoter for producing dicotyledonous and manuscityledonous transgenic plants comprises a nacicic acid sequence derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                 Sequence 207 HP; 67 A; 58 C; 21 G; 61 T; 0 other,
                                                                                                                                                                                                                                                                                                                                                                                                                                       expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a g box operably or functionally linked upstream of a CAAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents the chimeric promoter Mpr1098. The promoter is derived from the petE promoter from pea plastocyanin generand contains only the TATA and CAAT boxes, and corresponds to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Page 67; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-587667/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200056906-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         plastocyanin gene promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA96460 standard: DNA; 207 HP.
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                                                                                                                                                                                                                                                                                                                                                  species, e.g. wheat, barley, wat, rice, or corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            promoters of the invention. The specification describes a chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transgenic tobacco plants. The promoter is used to construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pe(E promoter; chimeric promoter; transgenic plant; MPr1097
     136
                                                                                         604 a88-cc88tcc8c8tctttatcaccc8ttctataaaaaat a.actttytyaytctaca. 662
                                    663 tittguttocottouaacacutusuuagugaagagactaattaattaat 710
                                                                                                                                                                                  544 ctctgtggcaratctacattatctadatcucucuttcttccucucutctgagccaracaa 603
                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                             76 aaacccaatgcacatotttatcatccattctataaaaaatraccttctqtqtqtctctct
ttogattocottoaaacacatacaaattoagtagagaagaaactoatt
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                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                        15.9%;
84.5%;
                                                                                                                                                                                                                    Score 116; DB 21;
Pred. No. 7.1e-14;
9; Mismatches 25;
                                                                                                                                                                                                                                                         Length 207;
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                                                                                                                                                    KESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Overy Match 14 08; Score 106.4; bs 21; beigth 280; Hest Local Similarity H8 08; Prod. No. 5.26 12; Matches 140; Conservative c Missoatches 16; beids 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The specification describes a chimeric expression promoter comprising a promoter of the pea plust expanin area, or comprising a 6 hox operably or functionally linked upstream of a CAAI hox. LAFA hox and transcription initiation site. The chimeric promoters are used in
AAA96.46.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transcription initiation site. The chimetic promoters are used in expression vectors for producing transgenic plants, such as directly reduced such as meaning species, e.g. potation, telephone, tenture, tomate, melon, encumber, pea, rape, bestroot, or sunflower, and monocotyledonous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            but not full light regulated expression in transgence tobacco plants. The promoter is used to construct chimeric promoters of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 promoter is derived from the petE promoter from per plastocyanin dene, by lusing the petE as 1 like and now enhanced like elements to the promoter McF1098 (comprising ERIA and CAAI boxes of petE), and then deleting 43 bp from the 5 mFR. The petE promoter directs cell specific deleting 43 bp from the 5 mFR. The petE promoter directs cell specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2: Page 69: B3pp. End.ish.
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                                                                AAA96461 standard; DNA; 281 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequences 250 HP, Bl A, E1 C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               species, e.g. wheat, burley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents the chimeric promoter Mpr1109. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              derived from a promoter of the pea plastocyanin dene
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plastocyanin dene promoter: s.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08 FEB 2001 (lirst entry)
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Rest Local Similarity 22.8%: Segre 89.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression promoter comprising a promoter of the pea plastocyanin ache, or comprising a 6-box operably or functionally linked upstream of a CAAI box. TATA box and transcription initiation site. The chimeric promoters are used to expression vectors to producing transactors plants, such as dicotyledonous species, e.g. potato, tobacco, centon, letture, tomato, melon, cucumber, pea, rape, bectroal, of sunflower, and monocutyledonous
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AAA96463;
                                                                              AAA96464 Standard; DNA; 280 BF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 281 Bb: 95 A; 71 C; 45 G; 80 T; C other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             species, e.g. wheat, burley, out, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      promoter directs cell specific but not full high regulated expression in
transgenic tobacco plants. The promoter is used to construct chimeric
promoters of the invention. The specification describes a chimeric
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 68; B3pp; Emglish
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ramer I. Sinber V.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28 SEP 2:00.
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                                                  RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operably or functionally linked upstream of a CAAT box, TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous species, eq potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, beetroot, or sunflower, and monocotyledonous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 promoter is derived from the pelE promoter from pea plastocyanin gene, by fusing the petE as-1 like and nos enhancer like elements to the promoter Mep1098 (comprising TaTa and CaAT boxes of petE), and then inserting a G box and fusing a fragmentof 35S RNA of CMV. The petE promoter directs cell specific but not full light-regulated expression transgenic tobacco plants. The promoter is used to enstruct chimeric promoters of the invention. The specification describes a chimeric promoters of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 2; Page 70; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleotide sequence of
                AAA96465 standard; DNA; 296 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plastocyanin gene promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 280 BP, 88 A; 73 C, 42 G, 77 T, 0 other.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            species, e g. wheat, barley, cat, rice, or corn.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rance I, Gruber V,
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                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                    11
                                                                                                                                                                                          promoter, chimeric promoter; transgenic plant; MP11110;
                                                                                                                                                                         teacettetgtgtgtetetetttegatteeetteaaaracatacaaattraytayayaay 247
                                                                                                                                                                                                                                              ccacacatcttagccacacaaaaacccaatccacatctttatcatccaffctafaaaaa
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                                                                                                                                                                                                                                                                                                                                                                                                       DB 21; Length 280
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric expression promoter for producing dicotyledonous ar
monoecutyledonous transgenic plants comprises a nucleic acid
derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for producing transgenic plants, such as diestyledomous species, e.g. petato, tobacco, cotton, lettuce, tomato, melon, eccumber, pea, rape, beetroot, or sunflower, and monocotyledomous species, e.g. wheat,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents the chimeric promoter Mprillil. The promoter is derived from the petE promoter from pea plastocyanin gene,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 296 BP; 94 A; 74 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               by fusing the petE as 1 like and nos enhancer like elements to the promoter Meria98 (compaising IAIA and CAAT boxes of petE), and then
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Pred. No. 1.2e
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Rest Local Similarity
                                                                                                                                                                                                                                                                                             Matches 144; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              construct chimeric promoters of the invention. The specification describes a chimeric expression promoter comprising a promoter of the postplastocyanin general constitution of box operably or functionally linked upstream of a CAAL box. TAIA box and transcription initiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chimeric expression promoter tor producing stootyledomous and monorotyledomous transferic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyunin deme
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 303 BP; 101 A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2: Page 7(; 83pp) English.
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Pred. No. 1.2e 08;
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IID AAA8
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                                                                                                                                                                                                                                                                   Query Match 12.1%; Shore 88.4; DR. West Local Similarity #1.0%; Pred. No. 1.60 #8 Matches 115; Conservative 0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                               Specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a 0-box operately on tunctionally linked upstream of a CAAL hox. JAIA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dioutybedonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pearape, beetroot, or sunflower, and monocoryledonous species, e.g. wheaf,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inserting a G box and tusing a tragment compressing a duplication of the element as 2 and as 1. A 72 by fragment compression the elements as 2, as and as 1 was then deleted, the petE promoter directs self-specific but not rull light regulated expression in transpense tobacco plants. The promoter is used to construct chimeric promoters of the invention. The promoter is used to construct chimeric promoters of the invention. The
                                                                                                                                                                                                                                                                                                                                                                                                                              barley, oat, rice, or corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents the chimeric promoter Mpill43. The promoter is derived from the pets promoter from per plastocyanin dense by Jusing the pets as I like and now enhancer like elements to the promoter MPI1098 (comprising TATA and CAAT boxes of pets), and then
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 72; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric expression promoter to producing discryledonous and monocutyledonous transquire plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyania dene
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 220 BP: 35 A; 6- 3: 26 G; 6: 1; p office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wr-z()uu5f 90f.A)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       petE promoter; chimeric promoter; transgenic plant; Mrill48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleotide sequence of the promoter MPr1143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 MAR - 2000; 2000Wo 1B00413
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RESULT 15

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AAT 17099
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence represents the pea plastocyanin promoter enhancer corresponding to nucleotides 444 to -179 of the full length sequence. This enhancer has been used to increase the expression of the GUS reporter gene in tobacco where the GUS gene was under the control of the PetE producter and 355 CAMV promoter the cuba control of the PetE producter and 355 CAMV promoter the cuba control of the PetE producter and 355 CAMV promoter the cuba control of the PetE producter.
                                                                                                                                                                                                                                                                                                                                                     a plant increases in green or non-green tissues, especially roots, tubers, seeds, stems, flowers or leaves. This enhancer may operate both in normal and reverse directions and can be attached either to
                                                                                                                                                                                                                                                                                                                                                                                                          also used joined to the \cdot 330 to \cdot 1 region of the patatin promoter PS20 which was joined to the GUS reporter and transformed into polato using A.tumefaciens. Expression of the gene promoter of one or more genes of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 9; Figure 6; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Enhancer from pea plastocyanin promoter region—used to produce transgenic plants having increased expression of a desired gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1997-310610/28.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29 NOV-1995,
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                                                                                                                                                                                                                                                                                                      Sequence 266 BP; 102 A; 47 C; 34 G; 83 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                        the promoter or terminator of the gene to be expressed.
223 teacageaalytteateagalaacreaettlaageceaegeact 266
                                 502 teacaacaateetqatgagataacceaetttaageeeaegeaet 545
                                                                                                                                                                          163 tagtattatttaetaaaaaaaatetaageeaegteggaggataaeateeaaeeeageeaa
                                                                                                                                         103 tettagattttataetteattgattattteatagageaagtaggagaaataaaaatatae 162
                                                                                                                                                                                                                                 y Match 10.8%; Score 79.2; DB 1 Local Similarity 67.7%; Pred No 9 7e-07
                                                                                      Sandhu JS,
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA96477;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents the chimeric promoter Mpr1112. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 25; Page 79; B3pp; English
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                                                                                                                                                                                                                                                                    Conservative
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Search completed: September (1, 1, 0.2, -0.1; 64, 22, 16b, 11mer, 8074, see

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Query Match Length DB	D#	Πυ	Description
	95 4	13 1	611	10	BI 271179	BI271179 NF0=0010F
2	85 8	11.7	683	16	BF637150	
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c 11	72.8	10.0	376	10	BF635678	
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B1645072 OP2862 M1	AL075432 Drocophil		A! 9=2544   Presophil	AL063632 Drosophil	AL189701 Tetraodon	AL097301 Drosophil	AL174271 Tetraodon		ALICESUB Tetraodon				ви001091 Т122545b	AL547503 AL547503		Anderson Jakiton		AL097768 Drosophil	AL427304 clone BA0	A1.054918 Drosophil	A#167066 SP_0094_8	AF 14901 AL514901	AL096962 Drosophil	AL066801 Drosophil	AL069526 Drosophil	BM358135 CAFa000	ALCOSORU DIOSOPILI

## ALIGNMENTS

	FEATURES Source	REFERENCE AUTHORS TITLE TOURNAL COMMENT	RESULT 1 BI271179 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM
/organism medicato truncatura" /db_xrof-"taxon_1880" /clone "NF650210FL" /clone_ib="De-veloping_ilowers" /tlasue_type="De-veloping_ilowers" /tlasue_type="De-veloping_ilowers" /dev_stage="De-veloping_ilowers" /dev_stage="De-veloping_ilowers" /dev_stage="De-veloping_ilowers_ilowers_and_flowers_ilowers_and_flowers_ilowers_and_flowers_ilowers_and_flowers_ilowers_and_flowers_ilowers_and_flowers_ilowers_and_flowers_a	The Samuel Roberts Noble Foundation 2510 Sam Noble Parkway, Ardmore, CK 73402, USA Tol: SRC 221 7361 Fax: 58C 221 7380 Email: qdmay9noble.org Insert Length. 611 Std Error: 0.00 Plate: 050 row: C column: 10 Sed primer, TCACACAGGAAACAGCTATGAC. Location/Qualifiers 1. 611 Authority Medicaro transatula*	l (bases 1 to 611) l (bases 1 to 611) l (bases 1 to 611) Fortus-Jeruz, L., Scott, A.E., Harris, A.E., Conzales, R.A., Beil, C.J., Felores, H.E., Imman, J.T., Weller, J.W., and May, G.D. Expressed Sequence Tags from the Samuel Roberts Noble Foundation Medicago truncatula flower library Inpublished (2001) Contact: May GD Contact: May GD	H1271179 H271179 H271179 H271179 NPO50C10F11F1082 beveloping flower Medicago truncatula cDNA clone B1271179 B1271179 B1271179 B1271179 B1271179 B1271179: G1:14879432 EST. Marrel medic. Medicago truncatula Eukaryota, Viridiplantae, Streptophyta; Babiyophyta; Trachcophyta; Spermatophyta: Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta: Magnoliophyta; Pabaceae; Papillonoideae; Trifoliege, Medicago: eurosids I; Fabales; Fabaceae; Papillonoideae; Trifoliege, Medicago

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Flores,H.R., Inman,J.J., Weller,L.W. and May,C.D.
Expressed Sequence Lys from the Samuel Experts Noble Foundation
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Spermatophyta; Madnollophyta; eudicotylodous; Pore eudicots;
Rosidae; eurosids I; Fabales; Fabareae; Papillonoldeae; Tritolneae;
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the Sammel Roberts Noble Featidation
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Unpublished (2000)
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clone NEIO5G07PL 57, mRNA sequence.
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a 148 c
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truncatula plants were transplanted to phosphate tree sand
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                                                                                                                                                                    392 GGCMATATAWAWWWWTTTTTTTAWAAAWAAATAATTWWAAWAWAAAAATTWWAAAA 451
                                     181 gaattittggcaagtcattaaaaagaaagaataaattattittaaaattaaaacttgagt 240
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                                                                                                                                                                                             please see http://www.fruitfly.org The BuGP prosophila mclanogaster BAC library was prepared by Kazuloyo Secegawa and Aaron Mammoser in Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI-9R and was constructed by partial EcoRI digestion of Decsophila DNA provided by the BuFP from the isogenic Strain v2. on bw sp. the safe strain used for the BUFP's and how to order individual BAC clones, the entire library, or library for the training from the BACPAC Resource Center can be found at the Acceptable from the BACPAC Resource.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Drosophila melanogaster genome survey sequence TET3 end of BAC # BACR08K10 of RPCI-98 library from Drosophila nelanogaster (fruit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     - Web . www.yenuscope.cns.f1)
Determination of this BAC-end sequence was carried out as part of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (02-JUN-1999) Genosoppo - Centre National do Sequencage BP 191 91006 EVRY cedex - FRANCE (E-mail : Segref@genoscope.cns.fr
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Pterygota, Neoptara, Endopterygota, Diptera; Brachycera;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organ sm-"Drosophila mclanogastcr"
/db_xref="taxon:7227"
/dlone_lib-"RPCI-98"
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a 64 c 131 g 202 t 503
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NEO14F06LF1F1058 Developing leaf Medicago truncatula
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Contact: May GD
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                                                                                                                                                                                                                               Seq primer: TCACACAGGAAACAGCTATGAC
                                                                                                                                                                                                                                                                                           Medicago Genome Initiative accession: MG1:8:14974 
Insert Length: 671 Std Error: 0.00
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The Sammel Roberts Noble Foundation
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/clone_lib-"Developing leaf"
/tissue_type="leaf"
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                                                                                                                                                                                        please see http://www.fruiffly.orq the BDCP brosophila metamonaster BA' liberry was prepared by Kazuloyo isoccawa and Arron Manmosen in Pieter de Jone's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPC' 9H and was constructed by partial Brokel didestion of Ecsophila DNA provided by the RBCP from the Isocence Stroin y2: the bw sp. the same strain used for the HDCP's Plant EST libraries. A more detailed description of the library and how to order individual HAC clones, the entire library, or litters for hybridization from the BACPAC Resource Center can be found at http://bei.sed.buffal.ed.
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The BDGP is constructing a physical map of the Drosophila
melanogaster genome using these BACs. For further information
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Contact : Pascal Gamas and Effenne Pasca; Former (Laborational)
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Xholf N. Cruncatula sterilised seeds were germinated for
22h at 26 S, leburg transplanting into a Tylegolisses skil
278 calcinod lernagheen mix in the presence of onion
roof tradments colonized by the arbuscular mycorrhical
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Zenftivar "Jemalong"
Zdb_aref: "taxon: 880"
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/elone_lib_"MtBC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP 191 91006 EVRY codex - France
Email: seqref@gencscope cns fr
Email: seqref@gencscope cns fr
Contact: pascal Gamas and Etienne-Pascal Journet, Laboratoire de
Biologie Moleculaire des Relations Plantes-Microorganismes,
CNRS-INPA, RP 27 31326 Castanet Tolosan Cedex, France (Email:
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MTHCTTHCOPT MCHC Modicago truncarula orna olono MthcTTHCOP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Medicago truncatula ESTs from endomycorrhizal roots Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Journet, E.P., Grespeau H., van-Junee, F., Gowy, F., Gaillon, C., Niebel, A., Carreau, V., Chatagnier, C., Kahn, D., Gianinazzi-Pearson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridipiantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae, eurosids I, Fabales; Fabaceae, Papilionoid-a-; Trifolieae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                http://sequence.fooloose.inra.fr/Mfroncatula html)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mt-est@toulouse.inra.fr Website
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genoscope - Centre National de Sequencage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact. Genoscope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   but with a high level of nitrate. After 3 weeks RNA was extracted from whole nort systems. FINA was propared from polyA+ enriched RNA. The cDNA was directionally ligated into Uni-zap XR vector from Stratagere and packaged using Gigapack Gold packaging extracts. Plasmids containing cDNA
root fragments colonized by the arbuscular mycorrhizal fungus Glomus intraradices (Schenck & Smith, isolate LPA8). The plants were watered every day and twice a week with
                                                                                             /note-"vector: pBluescript pSK; Site_1: EcoRI: Site_2: xhol; M. truncatula storilised seeds were germinated for 72h at 25 C. before transplanting into a 1/3 Epoisses soil : 2/3 calcined Terragreen mix in the presence of onion
                                                                                                                                                                                                                                                                                                                                                                                    /organism-"Medicago truncatula"
/cultivar-"Jemalong"
/db_xref "taxan 3880"
                                                                                                                                                                                                                                                     /tissue_type:"arbuscular mycorrhida"
/dev_stage="harvested 3 weeks post inoculation with Glomus
                                                                                                                                                                                                                                 intraradices"
                                                                                                                                                                                                                                                                                                                                                      /clone-"MtBCl1H02"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GSS; genome survey sequence. 
Tetraodon nigroviridis.
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Tetraodon nigroviridis genome survey sequence 17 end of clone
22211 of library 3 from Tetraodon nigroviridis, genomic survey
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Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformos;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tetraodon nigroviridis
                                                                                                                                                                                                                                                                                                                                                                                                                                                  http://www.genoscope.cns_fr/Tetraodon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (12-A) F-2500; to the LMH.
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Bouneau,L., Biliault,A., Quetier,F., Saurin,W., Nernot,A. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AL175696.1 GI:7813753
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roest-Crollius, H., Jaillon, D., Dasilva, C., Bouneau, L., Fisner, C.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inserts were mass-excised from phage stocks using FXAssit helper phage and propagated in SOLK cells. Clone ordering and sequencing was performed by the Centre National de Sequencage (Genoscope, Evry, France). Note: EST may be of lungal origin."
/clone_lib-"G"
/note-"Genescape sequence in : C@AC222CFARLETECND : 17"
/note-"Genescape sequence in : C@AC222CFARLETECND : 17"
                                                                                                                                                                                                 /close-"222Lll"
                                                                                                                                                                                                                                      /organise="Tetrapdon nigroviridis"
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ጵፍ ዓ*, Pred No. 0.0074;
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                                                                                                                                                                                          d'Etude du Polymorphi ame Humain) with funding provided by a MRC
project grant. The DNA was prepared from embryos by Alain Bucheton
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HP 191 91006 EVEX cedex - FRANCE (E mail | Segret genoscope.cos.tr
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/organism "Trosophīla melanepaster"
/plasmid "plocloBACII"
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                                                                                                        d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alair Buchcton and Genevieve Payan. It has been constructed in the vector

    Web: www.genoscope.cns.fr)
    Determination of this BAC-end sequence was carried out as part of a

                                                                                                                                                                                                                                                                                Submitted (23-JUL-1999) Genoscope - Centre National de Sequencaye
BP 191 91006 EVRY cedex - FRANCE (E-nail : seqref@genoscope.cns.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Drosophila melanogaster
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Insert Length 376 Std Error: 0.1
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Pterygota; Neoptera; Endopterygota, Diptera; Hiachycera;
Muscomorpha; Ephydroidea, Drosophilidae, Drosophila.
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/dev_stage="Pooled timepoints"
/note="Vector: Lambda Zap; Contains a mixture of entire
/note="Vector: Lambda Zap; Contains a mixture of entire
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/organism="Drosophila melanogaster"
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           BF 191 9100 EVRY cedex FRANCE (E mail sequelegenoscope.cns.fr )

Determination of this BAC end sequence was carried out as part of a collaboration with the Berkeley Drosophila Genome Project (BDGP). The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC library was prepared by Kazutoyo Osoegawa and Aaron Nammosor in Pieter de Jong's laboratory in the Department of
                                                                                                                                                                                                                                                                                                                                                                                                                                                       fly), genomic survey sequence.
AL063921
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Cancer Genetics at the Ruswell Park Cancer
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Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
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Local Similarity 43.6%; Pred. No. 0.02;
hes 198; Conservative 21; Mismatches 235; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         collaboration with the Berkeley Drosophila Genome Project (BDGP). The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACS. For further information please see http://www.fruitfly.org.The BDGP @rosophila melanogaster BAC library was prepared by Kazutoyo Gsoegawa and Aaron Mammoser in Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI 98 and was constructed by partial EcoRI diges ion of Drosophila DNA provided by the BUGP from the isogenic strain y2; on bw sp. the same strain used for the BUGP's Pl and EST libraries. A more detailed description of the BUGP's and how to order individual BAC clones, the entire library, or filters for hybridization from the HACPAC Resource Center can be location/Qualifiers.
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    Web: www.genoscope.cns.fr)
    Determination of this BAC-end sequence was carried out as part of a

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Eukaryota, Metazoa, Arthropoda, Tranheala, Hekapoda; Insecta;
Plerygota, Neoptera; Endopterydota; Dipleza; Brachycera;
Muscomorpha, Ephydroidea; Drosophilidae, Drosophila.
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## ALIGNMENTS

ALLOW NUMBERICATION DATE: PEICATION DATE: PEICATION DATE: NUMBERICATION	WESULI 1  US-08-232-463-14/c  Sequence 14. Application US/98222463  Patent No. 5670387  GENERAL INFORMATION: APPLICANT: DORNER, F. APPLICANT: SCHEIFLINGER, F. APPLICANT: APPLICATION DATA:  USA ZIP: 22313-029  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARP: Patentin Pcicasc #1.0, Versicr #1.25 CUKKENT APPLICATION DATA:
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                                                                                                                                       TELEFAX: (414) 271 652 INFORMATION FOR SEQ 10 No.
                 MOTHSTRUM
                                                                                                                                                                                               REFERENCE, TRACKET NUMBER OF
                                                                                                                                                                                                                                                                 CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: Baker, Jean C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              THILE OF INVENTION: FIRER SPECIFIC PROMOTERS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: THM PC compatible
OPERATING SYSTEM: PC bos/MS bos
                                     PoPOLOGY: linear
                                                                                TYPE
                                                                                                                                                                                                                                                                                                                                FILING DATE:
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                                                      STRANDEDNESS:
                                                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                                                                             REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                                                                               LENGTH:
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bucteric acid
DEDNESS: double
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               fortunents) VNc.
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US-08-467-504 4/c
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: MOLECULE LYPE: DNA (demonic) 0.08-0.8-46.7-5.04-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: John,
                                                                                                                        TELEPHONE: (414) 277-5709
TELEPAX: (414) 277-5709
TELEPAX: (414) 271-552
INPORMALION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PT DOS/MS DOS
SOFTWAKE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: EDLATE PREMOTER
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                                                            STRANDEDNESS:
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                                         : Atmorphed
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                                                                                                       LENGTH:
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                                                                                                     1974 base pairs
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411 East Wisconsin Average
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Query Match 6.6%; Score 48; DB 4; Longth 974; Host Local Similarity 50.4%; Pred. No. 0.036; Matches 143; Conservative 0; Mismatches 140; Lidels

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; ORGANISM:
US-)8-998-416-288
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US-(8-998-416-288
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                                     TOPOLOGY: line
MOLECULE TYPE: I
ORIGINAL SOURCE:
                                                                                                                                                                                      NFORMATION FOR SEQ ID NO: Z88:
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APPLICANT:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541 8587
                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: CH 0016/97 FILING DATE: 31-DEC-1996 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Wendland, Jurgen
APPLICANT: Knechtle, Philipp
APPLICANT: Rebischung, Corinne
APPLICANT: Rebischung, Corinne
TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHRYA GOSSYPII
TITLE OF INVENTION: AND USES THEREOF
NUMBER OF SEQUENCES: 1152
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                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                        TELEPHONE. 919-541-8689
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER FILING DATE: 24-DE CLASSIFICATION: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                 LENGTH: 837 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                            NAME: Meigs, J. Timothy PEGISTPATION NUMBER 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE:
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Research Triangle Park
: No. 6239264th Carolina
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Steiner, Sabine
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                      PAG1241RP
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                                                                                  linear
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                                                              DNA (genomic)
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Query Match

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Score 47.4;

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Matches 165; Conservative
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                                                                                                                    TELEPHONE: (619) 235-8550
TELEFAX: (619) 235-0176
INFORMATION FOR SEQ ID NO: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IHM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0. Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
HYPOTHETICAL:
                MOLECULE TYPE:
                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                         REFERENCE/DOCKET NUMBER: NIH121.001CP1 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                            AFFORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           380 TAAATTATTATTTATTATAATATCTATTTTTATAAATATCTGTTGATTTATATATTAT 439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   320 TAATTGTTTATTAAAATAATATTATTATTATTATTAAAAGATTTAATTTATTTAATATTO 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  130 taattgetgtaaataaataaggatgaegeattagagagatgtaeeattagagaattittig 189
                                                                                                                                                                                                                                                                 FILING DATE: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 620 Newport CITY: Newport Beach
                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE:
                               : ADGNODOL
                                                                                                                                                                                                            NAME: Israelsen, Ne REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER
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                                                                                    LENGTH:
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                                                                   nucleic acid
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                                                                                    19124 base pairs
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Miller, Louis H
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                                 linear
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N: 435
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0; Mismatches 176;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/0786 106 Patent No. 5389526
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Obery Match 6.8%; Score 46; DB 2 Longth 19124; Best Local Similarity 4.6%; Pred. No. 0.12; Mismatches 20%; Conservative c. Mismatches 30%; Endels 0. Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1562? ΑΛΑΛΑΤΩΑΤΤΑΊΑΛΑΛΑΛΑΛΑΙΑΛΩΛΑΩΑΛΩΑΛΩΑΛΑΛΑΛΑΛΑΛΑΛΙΩΑΤΤΑΛΑΛΑΛΑΛΑΛΑ
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                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Slade, Martin B
APPLICANT: Chang, Andy C M
APPLICANT: Williams, Keith:
                           ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Improved Plasmid Vectors for Cellular
TITLE OF INVENTION: Shime Moulds of the Genes Dictyostelium
NUMBER OF SEQUENCES: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ALBRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               | 181 qaatttttqqcaaqtcattaaaaqaaqtaaattatttttaaaaqttaaaqttqaqt 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               121 aataagggttaattgrigisuutsuataaggatgacgcultagagagatgtaccaltaga 180
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                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: DO HOSZMS DOS
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                                                      FILING DATE:
                                                                         APPLICATION NUMBER: DOL/A: 90/00530
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                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                      STATE
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One Liberty Place 46th Floor
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Joanne Long
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6.8%; Score 45.8; DB ]; Death 5852;

Best local Similarity 47.8%; Pred. No. 0.11;

Matches 164; Conservative 0; Mismatches 177; Indels 1; Gaps
                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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                                                                      COMPUTER READABLE FORM
                                                                                                                                                                                                                 APPLICANT: Gruidt, Michael E.
TITLE OF INVENTION: No. 5721852cl Entonopoxvirus Expression System
NUMBER OF SEQUENCES: 40
                                                                                                                                                                                                                                                          APPLICANE: Moyer, Richard W. APPLICANE: Hall, Richard L. APPLICANE: Gruidl, Michael F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
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TELECOMMUNICATION INFORMATION:
TELECHONE: 215 568 (100
                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   129 ttaattgetgtabatbabtbabggdtgaegeattagagagatgtaeeattagagagaittitt 188
               MEDIUM TYPE, Floppy disk
COMPUTER: THM PC compatible
COPERATING SYSTEM: PC 18-82/MS-18-8
                                                                                                                               STREET: 2+2.
CTTY: Cainesville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              255 TATLITTITITITCAAACITAAAAAGATAATITITITTITAATA STIGATGACTIT 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  249 taaanatgigattatiituuigaaiigaalgaaga vagiigaatiaaagiigtaliagiaai 307
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                                                                                           21P:
                                                                                                                                          ACHRESSEE: David B. Sallmann. Suite A. STREDE: 2421 M.W. 41st Street, Suite A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           468 ajtouditaaotoatititatutitoatugafoudutuagagagaadaacggglutultudto 427
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                                                                                                           COUNTRY:
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                                                                                                                                                                                                                                                                                                                                    i, Application Highwelt7755A
o. 5721352
                                                                                    32606
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                                                                                                                             Florida
                                                                                                         U.S.A.
Patentin Release #1.0, Version #1.2
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/01 FILING DATE: 19-AUG-1993

US/08/107,755A

CLASSIFICATION:

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Best Local Similarity
Matches 156, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (904) 372-5800 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 07/657,584
FILING DATE: 19-FEB-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/827,658
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                       2135 AAAAGTGTTCCTACGTTGATATTTAAAAAATGATATTATAGAGTTATATTTATCTA1TTAAT 2076
                                                                                                                                                                                                                                     2194 TATTCATTTAATAAAAAAAATATGTTAGTGATTATTACTAAA-TATTATGGATTTATGACA 2136
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                                            2015 ACGGGTATTTCTCCCTATAATTACATATTTATTATATCTAGATTTCATAAATTCTATATGT 1956
                                                                                                          2075 CTCACATTAGTTATAATACCAAAATTATTTAATATATATTATTAATAGATATATTATTTT 2016
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408 gaaataacggtatattaat.ccct 430
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REFERENCE/DOCKET NUMBER: UF114.C2
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                                                                          cetatatattgeeeeataeagteagttaaeteatttttatattteatagateaaataaga 407
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nucleic acid
DEDNESS: double
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1474..2151
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3080..6091
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2502..2987
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                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                              Score 45.4; DH
Pred. No. 0.14;
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                                                                                                                                                                                                                                                                                                                                                                                            DH 1; Length 6768;
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RESULT 8
US-07-991-867B-1/c
US-07-991-867B-1
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Patent No. 547678
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APPLICANT: Hall, Richard L.
APPLICANT: Grudf, Michael E.
TITLE OF INVENTION: No. 5476781el Entomopoxvirus Expression System
                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 8457 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 19-FEB-1991
ATTORNEY/AGENT INFORMATION:
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SOFTWARE: Patentin Release #1.0, Version #1
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REFERENCE/DOCKET NUMBER: UF114.C3
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TELEFAX: 904-372-5800
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KEGISTRATION NUMBER: 31,79
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IBM PC compatible
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              complement (6277..6768)
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ATTORNEY/AGENT INFORMATION
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                     APPLICATION NUMBER: US 07/657,584 FILING DATE: 19 FEB 1991
                                                                                          FILING DATE
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                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
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computer: IBM procompatible
operating system: probosyms has
software: Patentin Release #1.0, Version #1.25
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2421 N.W. 41st Street, Suite A:1
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Hall, Richard L.
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US-08-487-826B-13/c

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. Sequence 13, Application US/08487826B : Patent No. 5993827

GENERAL INFORMATION:

APPLICANT:
APPLICANT:

Sim, Kim L. Chitois, Chetan Miller, Louis H

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1955 GATACIATATCIAATACATCACI 1933
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Overy Match 6.2%; Score 45; DB 2; Length 19124; Hest Local Similarity 45.1%; Pred. No. 0.2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (619) 235-8550
TELEFAX: (619) 235-0176
NFORMATION FOR SEO ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 19124 hasc
                                                                                                                                                                                                                                                                                                                                                                                                         18301 TTGTTTGATAAAAGAATTAAGAAATTAAAAAAGGAATAAAGTTAAATAAAAATAAGAAATT 18242
                                                                         18357 TTTTTT----TTTTTTTTTTTTTTTTTTTTTTGTTGATATGTAGATTACTTTATGTTTT 18302
                                                                                                                                                          18477 TCTTAACTCTTTCTATGTTAAAATTATTATTATAATCTCAATATGTTTTAATCGTAATA 18418
                                                                                                                                                                                                                                                                                                                         18657 AAGACCTATATAAGTATASAAACGACATAAAATATATCGATAAAATCGAAAATTATSTINA 18598
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
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                                                                                                               326 ttaatttgacatttgatc_tttcctatatattgccccatagagtcagttaactcattttt 385
                                                                                                                                                                                                 266 aatgaattgatgaaagag"tggattaaagttgtattagtaattagaatttggtgtdaaat 325
                                                                                                                                                                                                                                                                              206 aaagaataaattattittaaaaltaaaagttgagtcatttgaltaaacatgtgattattt 265
                                                                                                                                                                                                                                                                                                                                                                146 ataaggatgacgcattagagagatgtaccattagagaattttttggcaagtcattaaaaag 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                86 caaatatcattgaggaathtgacaaaagctacacaaataagggttaattgctgtaaataa 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 10-SE
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                                    92660
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California
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Wellems, Thomas E.
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                                                                                                                                                                                                                  Matches 115;
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INFORMATION FOR SEQ ID NO:
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NAME: Melds, J. Timothy
REGISTRATION NUMBER: 38,241
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                               ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 837 base pairs
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303 gtaattagaalttggtglcaaalttaatttgacalttgalctttlcctalalalattgcccc 362
                                                                                                                                                                 183 attitiggcaagtcaitaaaaaggaaggaataaaftaffittiaaaattaaaagfiiggtcai 242
                                                                     243 tttgattaaacatgtgattatttaatgaattgatgaaagagttggattaaagttgtatta 302
                                                                                                                                            CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Fluppy disk
COMPUTER: IBM PC compatible
OPERATING SYSIEM: PC-POS/MS-POS
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STATE: No. 6239264th Carolina
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/INDOKET NUMBER: PE/5-30306/A/CGC1976
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Steiner, Sabine
Mohr, Christine
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Knechtle, Philipp
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                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                              DNA (genomic)
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US 08-991-677-]]
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                                                                                                   Query Match 6.1%; Score 44.6; DE 6; Length 50%;
Hest Local Similarity 46.8%; Fred. N., 0.18;
Matches 210; Conservation on Mismarches 234; Indels of
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APPLICANT: Smeltzer, Richard H
TITLE OF INVENTION: Production of Syringy! District to Cymnosperms
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                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 19
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ST, JENS, RIENECK, KLAUS, WIND, ANNETTE; DAKORSEN, PALLE
2930 AAAAAAAAATATATGAATTITA: TAGCTAAGAATATAAATCAGAACATAGATTTATTGT 2871
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                                            24. ataaaaacataatidtaaqtataa adaaaadtidtacaadagtidtaccaaaatagtid 83
                                                                                                                                                                                                                                                                                                                    FILING DATE: 18 SEP 1583
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                                                                                                                                                                                                                                                                  LENGTH: 1095
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                                               TELLEAK: 919-541 8589
INFORMATION FOR SEQ ID NO: 186:
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APPLICANT:
                                                                                                                                                                                                                                      PRIOR APPLICATION DATA.
APPLICATION NUMBER: CH 0016/97
FILING DATE: 31 DEC 1996
SEQUENCE CHARACTERISTICS:
LENGTH: 615 base pairs
                                                                                                                                                                                                                                                                                                                                                                         SOFTWAKE: PatentIn Release #1.0, Version #1 COURRENT APPLICATION DATA:
                                                                                           TELECOMMUNICATION INFORMATION: 919-541-8587
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TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHBYA (PSSYFII
TITLE OF INVENTION: AND USES THEREOF
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CEERATING SYCTIM PC 1505/MS For
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STATE: No. 6239264th Carolina
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                                                                                                                                                                 NAME: Meigs, J. Time REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
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Knechtle, Philipp
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                                                    INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
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APPLICANT: Stewart, Thomas .

APPLICANT: Flores, Maria V

APPLICANT: O'Sulivan william J

APPLICANT: O'Sulivan william J

TITLE OF INVENTION: Nucleotide sequence encoding carbamoyl

TOTAL OF INVENTION: plosphate synthetase !!
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ORIGINAL SOURCE:
                                                                                                           TELEPHONE: 703-816-4000
                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.24 CURRENT APPLICATION DATA:
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                                                                                                                                            NAME: Mitchard, Leonard C
REGISTRATION NUMBER 29.009
REFERENCE/DOCKET NUMBER: 47-
                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/446,855A FILING DATE: 06-Jul-1995 CLASSIFICATION: 435
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ZIP: 22201-4714
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TOPULUGY: linear
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                                                                                         TELEFAX:
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STRANDEDNESS:
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EDNESS: single
                 nucleic acid
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                                   8920 base pairs
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IBM PC compatible
SYSTEM: FC-DOS/MS-DOS
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Best Local Similarity 47.6%;
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Query
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Gapop 60 0 , Gapext 60 0
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All Art " "AAA 11 2640"

Anote "Seminance to be used as a bromoter for respilation expression"
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                                                                                                                                            artificial sequence.

1 (bases 1 to 731)

Vezina, L.P. and D. Aoust, M.A.
                                                                                                                                                                                                  synthetic construct
                                                                                                                                                                                                                 synthetic construct.
                                                                                                                                                                                                                                                                            Sequence 3 from Patent W00125455
                                                                                                                    Promoter for regulating expression of foreign Patent: WO 0125455-A 3 12-APR-2001,
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expression
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                          /organism="synthetic construct"
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          artificial sequence.
1 (bases 1 to 326)
Rance, I., Theisen, M.
                                                                                                   Sequence 3 from Patent FR2791358
AX036597
                                                   synthetic construct
                                                                synthetic construct
                                                                                          AX036597.1 GI:11225192
 Patent: FP 2791458
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M. and Gruber,V.
Α 3 29-SEP-2000;
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Best Local Similarity 1:00% Prod. Bo. 8.46 B
Matches 57: Conservative o: Mismatches o
                                                              ORCANISM
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Hest Loral Similarity 100.0% Pred. No. 7.96 18;
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All is a plant nuclear proton binding to the segmence ANIATIVIDATE in the promoters of light regulated agrees (N. Gatta & A.R. Cashmore 'Binding of a Pea nactear protein to promoters of certain photographated genes is modulated by phosphory.ation.' The Plant CCLL 1, 1069-1077, 1989).
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481, .41°
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/cultivar-"Poitham First"
/db_xrof "taxon: 8888"
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/note-"G box bosology"
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/db_wist "taxon:32630"
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/product="plastocyanin"
/protein_id="CAA48393.!"
/db_xref="G1:4379383"
/translation="MA"
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                 /note="imperfect direct repeat 2"
134, .171
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39. .41
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/note="imperfect direct repeat
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21. .50
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/clone_lib="lambda EMBL3"
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synthetic construct
                                                            Sequence 6 from Patent FR2791358
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Rance, I., Theisen, M. and Gruber, V.
Patent: FR 2791358-A 13 29-SEP-2000;
                                               AX046600.1 GI:11226195
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813. .1319
/note-"pracursor peptide (AA -69 to 99)"
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/note="Directional building block S3-Directional building
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/translation-"MATVTSTTVATPSESGLKTNAATKVSAMAKIPTSTSQSPRICVR
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100.0%; Pred. No. 5.9÷-06
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artificial sequence
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promoter MpT2098"
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Matches 41: Conservative 0; Mismatches 0: Indels
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synthetic construct
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                  synthetic construct
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d 24 e 45 d
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position of MPr1098, an 18 bp element equation of the element
and lusing a sequence of 98 bp (Suplication of the element
as 2 and as 1)
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1 (bases 1 to 303)
Rance,I., Theisen M. and Gruber,V.
Patent: FR 2791358-A 8 29-SEP-2000;
MERISTEM THERAPEUTICS (FR)
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Rance, I., Theisen, M. and Gruber, V.
Patent: FR 2791358-A 15 29-SEP-2000;
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Location/Qualitiers
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	Nevel premoter for regulating expression of foreign gener in transgenic organisms, more specifically in a leaf specific manner in transgenic plants.	WP1, 2001 2663:6727.		Venina L, D'Aoust M;	(MEDI:) MEDICAGO INC.		04-001-1999; 99US-0157129.	02-OCT-2000; 2000WO-CA01144.		100c-gag001	wuz00125455-A2.		Synthetic.	Promotor, transgenio plant, menos (ploden, like), letan epamipermo ds.		Promoter #1 for regulating expression of foreign genes.	to our wood (title energy)	ix (III - CO)) 7 (first cotty)	AASU2126;		AASU2126 Standard; DNA; 1350 BP.	AAS02126	JET 1	ALIGNMENTS	

Result		Query Match Length DB ID	Length	DB	ID	Description
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The sequence represents the coding sequence of promoter  $\pi_2$  for regulating expression of foreign genes in transquaric organisms. The promoter is

Claim 1; Page 9; 9pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 9; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel promoter for regulating expression of foreign genes in transgenic organisms, more specifically in a leat-specific manner in transgenic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MEDI-) MEDICAGO INC
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                             429 gaattittggcaagtcattaaaaagaaagaataaattattttaaaaattaaaagttgagt 488
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181 gaatttttggcaagtcattaaaaagaaagaataaattattttaaaattaaattaattagagt
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                 Similarity
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                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                      Score 731; DB 22;
; Pred. No. 5.7e-311;
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Claim 2, Page 67: 8 Spp: andi
                                                                         derived from a promoter
                                                                                                              monocotyledonous transgenic clants comprises a nucleic acid sequence
                                                                                                                                                   Chimeric expression promoter tor producing dicotyle lonous and
                                                                                                                                                                                                                                     WPT: 2000 587567/55
                                                                                                                                                                                                                                                                                                           Rance I, Gruber V.
                                                                                                                                                                                                                                                                                                                                                                                             (MERI ) MERISTEM THERAPENTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         petE promoter; elumente promiter; fransgemie plant; Merlüß6
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Best Local Similarity 100.0%; Pred. No. 4.3
Matches 57, Conservative 0, Mismatches
The present sequence represents the chumerre producter Myellow. The productor is derived from the policy and term for plantocyanin demonstration in 5° of the repeated invert sequences as well as the as I like box borne on a Sphi Traument of Alz by. The policy pounder directs cell specific but not full light regulated expression in
                                                                                                                                                            Claim 2: Page 66; 83pp: English
                                                                                                                                                                                                         derived from a promoter of the pea plastocyanin dene
                                                                                                                                                                                                                                   Chimerus expression promotes to producted discreptions and monocotypedomous transpense points comprises a naction and sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 326 Ber 111 At 91 Pr 48 0; 86 tr or other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        barley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rape, beetroot, or subliquer, and monocotyledonous species, e.s. wheat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        site. The chimetre promoters are used in expression vectors for producing transgenic plants, such as dicotyledomous species, each potato, todacco, cotton, lettoce, tomato, medeo, ejeumber,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operably of functionally linked upstream of a CAAT box. DATA box and transcription initiation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   promoter is derived from the petE promoter from peo plasfocyatin dene,
by a deletion in 5' of the repeated invert sequences, of the as 1 like
and enhancer like elements borne by two Sphl frauments of 403 bp. The
                                                                                                                                                                                                                                                                                                                                                                                                                  (MERL ) MERISIEM THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 MAR 2000; 2000WO 1B00317
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08 FEB: 2005] (first outry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA96458 standard; DNA; 623 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              petE promoter directs ceil specific but not full high regulared
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             110 agat aaree aett taagee earge acte tylygear aletaeattatet aaateaea 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     767 agataan caet traasi eranga ditetgiya arabetarat tatetaaateara 1823
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Pred. No. 3.3e 15;
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transaemic tobacco plants. The promoter is used to capetrat chime promoters of the invention. The specification describes a chameric

hans't in't inhimer in

expression promoter comprising a promoter of the pea plastocyanin

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RESULT
AAA96457
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   Matches
                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                        Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        are used in expression vectors for producing transgenic plants, such as dicotyledonous species, a gipotato, tobacco, cotton, lettine, tomato, melon, cucumber, pea, rage, beetroot, or sunflower, and monocoty.edonous
                                                                                                                                            expression vectors for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, bectroot, or sunflower, and monocotyledonous
                                                                                                                                                                                                                      promoter of the pea plastocyanin gene, or comprising a G hox operably or functionally linked upstream of a CAAT box, TATA box and
                                                                                                                                                                                                                                                                                                                  The present sequence represents the petE promoter from the pea plastocyanin gene. The fromoter directs cell-specific but not full
                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 65; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      species, e.g. wheat, barley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or comprising a 6 box operably or functionally linked upstream of a CANI box, TATA box and transcription initiation site. The chimeric promoters
                                                                                         Sequence 834 BP; 289 A; 175 C; 133 G; 237 T; 0 other;
                                                                                                                              species, e.g. wheat, barley, nat, rice, or corn
                                                                                                                                                                                                       transcription initiation site. The chimeric promoters are used in
                                                                                                                                                                                                                                                         specification describes a chimeric expression promoter comprising a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA96457 standard; DNA; 834 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 623 BP; 230 A: 128 C: 82 G: 183 T: 0 other;
                                                                                                                                                                                                                                                                             light-regulated expression in transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  767 agataacccactttaagcccacgcactctgtggcacatctacattatctacattatcaca 823
y Match
Local Similarity 100.0%; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gruber V,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimeric promoter; transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.28; Sc.
/ 100.08; Pr
                   Score 57; DB 21;
Pred. No. 3.1e-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 57;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
Mismatches
   3.le-15;
ກ່ອງ ດຸ
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3.2e-15;
                                     Length 834;
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Garis

Nucleotide Sequence of the promoter MPr1109

08 FEB 2001

(first entry)

AAA96452

AAA96462 standard; DNA; 250

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RESULT
                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             which is used to construct chimeric promoters of the invention in PCR reactions. The specification describes a chimeric expression promoter comprising a petE promoter of the pea plastocyanin genc, or comprising a G box operably or functionally linked upstream of a CAAT box, TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgence plants, such as dicotyledancus species, e.g. polato, tomato, nelon, cucumber, pea, rape, beetroot, or sunflower, and monocotyledonous species, e.g. wheat, barley, oat, rice, or corn, and monocotyledonous species, e.g. wheat, barley, oat, rice, or corn,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 24; Page 73; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The month of the second of the second of the second second
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                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 68 BP; 22 A; 19 C; 5 C; 22 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents the directional building block
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       derived from a promoter of the pea plastocyanin dene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chimeric expression promoter for producing directyledenous and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZERMAR JEEG, JOHNWO-IBO0317
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA96468 standard; DNA; 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EZ-MAK-JAHA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   790 cantotytyycacatotacattaintaaatoaca 823
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                             Score 34;
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                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                 DB 21;
                                                                                                                                                                                                                                                                                                             Length 68
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Hest Local Similarity 100,0%; Someoryative (c)
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W02000355305 AT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression vectors for producing transpenic plants, such as dicoryledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, par, cape, beetroof, or sunflower, and monocotyledonous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    a promoter of the per plaste yamin dene, of comprising a G box operably or functionally finked spatism of a CAAI box, IAIA box and transcription instration sate. The chimeric promoters are used in expression executes to:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    deleting 33 bp from the 57 BTR. The petE promoter directs cell specific but not full light regulated expression in transquare tobacco plants. The promoter is used to construct chimeric promoters of the invention.
                                                                                                                                                                                                             Pisum sativum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plasforyanin dene promoter:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08 FEB 2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         promoter is derived from the potE promoter from poliplastocyania gene
by fusing the petE as I like and nos enhancer like elements to the
promoter MP(1098 (compressing ATA) and CAAT boxes of petE), and then
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2: Page 69, B3pp; Englesh.
                                                                                                                                                                                                                                                                                                              Synthet in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        petE promoter: chimeric promoter: transdenic plant; MPr1097;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleotide sequence of the promoter MPr1098
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Best Local Similarity 100.0%: Fred. No. 3 years
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or comprising a 6-box operably or functionally linked upstream of a CAAI box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors by producing transcenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, exembler, pea, rape, bectroof, or sunflower, and monoccryledonous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents the chimeric promoter MgrIsbs. The promoter is derived from the pet E-promoter from pea plastocyanin demand contains only the TATA and CAAL bases, and contraresponds to a minimal reference promoter on the polE-promoter. The petE-promoter directs cell specific but not full light regulated expression in
Rance i, draber V.
                                      (MERI ) MERISTEM THERAFEUTICS
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                                                                              22 MAR 1999;
                                                                                                                      20 MAR 2000; 2000Wo [B00317
                                                                                                                                                              28 SEP 2000.
                                                                                                                                                                                                                                            Pisum sativum
                                                                                                                                                                                                                                                                                                      plastocyanin gene promoter;
                                                                                                                                                                                                                                                                                                                                                               Nucleotide sequence of the promoter MPr1110.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA96464 Standard; DNA; 280 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         are used in expression vectors for producing transgence plants, such as dicotyledonous species, e.g. potato, tobacco, contron, introp. Tomato, melon, cumber, pea, rape, beetroot, or sunflower, and monocotyledonous species, e.g. wheat, barley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expression promoter comprising a promoter of the pea plastocyanic gene or comprising a G box operably or functionally linked upstream of a CAAT box. TATA box and transcription initiation site. The chimeric promoters
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                                Claim 2; Fage 71; 83pp; English.
                                                                                     Chimeric expression promoter for producing dicotyledonous monocotyledonous transgenic plants comprises a nucleic ac
                                                                                                                                                                                                                                                                                        20-MAR-2000: 2000WO TROCP17
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                                                                  derived from a promoter
                                                                                                                                           WPI; 2000-587667/55.
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                                                                of the pea plastocyanin gene
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                                                                                  comprises a nucleic acid sequence
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0 00085;
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The present sequence represents the chimeric promoter Mpr1111.

site. The chimeric promoters are used in expression vectors for producing

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                                                                              The present sequence represents the chimeric promoter Mpr1153. The promoter is derived from the petE promoter from pea plastocyanin dence. The petE promoter directs cell-specific but not full light-regulated expression in transpence tobacco plants. The promoter is used to construct chimeric promoters of the invention. The specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inserting a 5 box and fusing a fragment comprising a duplication of the element as2 and as1. The petE promoter directs cell-specific but not full light-regulated expression in transgenic tobacco plants. The
describes a chimeric expression promoter comprising a promoter of the pea plastucyanin gene, or comprising a 5-box operably or functionally linked upstream of a CAAT box. TATA box and transcription initiation
                                                                                                                                                                                                                                                    Claim 2; Page 70; 83pp; English.
                                                                                                                                                                                                                                                                                                     Chimeric expression promoter for producing discipledonous monocotyledonous transgenic plants comprises a mack is an derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rance 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleotide sequence of the promoter MPr1153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA96464 standard; DNA; 303 HP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 296 BP: 94 A; 74 C; 45 G; 83 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for producing transgenic plants, such as dicotyledonous species, e.g. petalo, tobacco, cotton, lettuce, tomato, melon, cucumber,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanic gene, or comprising a 6 kex operably or functionally linked mystream of a CANT box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              promoter is used to construct chimeric promoters of the invention. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-MAR-2000; 2000WO-1800317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pisum sativum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plastocyanin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    petE promoter; chimeric promoter; transqenic plant; MPrll53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          barley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           790 cactetgtggcacatetacattatetaaate 820
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              beettoot, of sunflower, and monocotyledonous species, e.g. wheat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity 150.0%; Pred. No. 0.0
31; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene promoter; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DH 21;
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                                                                                                                                                                      Matches 2), Somery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Querry Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   promoter comprising a peter pointer of the per plastocyanin dene, or comprising a 6 box speciably of functionally linked apstream of a CAAT box. TATA box and transcription initiation show. Tata characteristics are used in express teams of the contract promoters are used in express teams species, e.g. polato, tobacco, contropolatos, such as dicotyledomous species, e.g. polato, tobacco, contropolato teture, tomato, melon, such as express e.g. wheat, barrey, out, fice, or corn, other and monocotyledomous species e.g. wheat, barrey, out, fice, or corn, other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chimeric expression proseder to producing dicotyledonous and monocotyledonous transquare plants comprises a nuclear acid sequence derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             transpente plants, such as directyledonous species, e.g. polato, tobacco, cotton, lettuce, tomato, melan, encumber, per, raps, beetroot, or sumflower, and monocotyledosous species, e.g. whear, barley, oat, rice,
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 72 BP; 22 A; 20 C; 10 G; 20 T; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 24; Page 75, # (pp: English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 303 BP; [0] A; [783].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR reactions. The specification describes a chimeric expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents the directional building block S5,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000 587667/55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               petE promoter; chimeric promoter; transgenic plant; MPr1108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         which is used to construct chameric promoters of the invention in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 MAR 1999; 99FR 000 0.45
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                                                                                   794 etgliggeacateraeallaretaaale 826
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    THE THE HELLIGHT AND TO A STATE OF THE STATE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                          2 (K. Shore 27) (DB 21) (Amath 72) (dd (DK) (Pred. No. 0.052)
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                                                                                                                                                                      Mismatches
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AAL/7099
weeky match
Best Local Similarity 100.0%; Score 27; DB-1H; Lendth 266;
Matches 27; Conservative or Mission
                                                                                                                                                                                                                                                                                        a plant increases in green or non-green tissues, especially roots, tubers, seeds, stems, tlowers or leaves, this enhancer may operate both in normal and reverse directions and can be attached either to
                                                                                                                                                                                                                                                                                                                                                                                             sequence. This enhancer has been used to increase the expression of the GUS reporter gone in tobacco where the GUS gone was under the control of the PetE promoter and 358 GAMV promoter. The enhancer was also used joined to the -300 to \pm 1 region of the parath, promoter PSZo which was joined to the GUS reporter and transformed into polaro asing Altumotaciens. Expression of the gene promoter of one of more dense of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence represents the pea plastocyanin promoter enhancer corresponding to nucleotides 444 to 129 of the full deagth sequence. This enhancer has been used to increase the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enhancer from pea plasfocyamin promoter region—used to productions genic plants having increased expression of a desired gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               promoter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pisum sativam plastocyanin promoter region
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                                                                                                                                                                                       Sequence 266 BP; 102 A; 47 %; 34 %; 83 I; 0 other
                                                                                                                                                                                                                                                          the promoter or terminator of the gene to be expressed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ADTE:) ADVANCED LECHNOLOGIES CAMBRIDGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29 NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26 NOV 1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 JAN:1998 (first entry)
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                                                                                                                                                                             Query Match 1.9%; Score 25; DB 21; Length 220; Best Local Similarity 100.0%; Pred. No. 0.37; Mitches 25; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents the chimeric promoter Mpr1143. The promoter is derived from the petE promoter from pea plastocyanin gene, by fusing the petE as-1 like and nos enhancer like elements to the promoter Mpr1098 (comprising TATA and CAAT boxes of petE), and then inserting a G box and lusing a fragment comprising a duplication of the element as2 and as1. A 72 br fragment comprising the elements as-2, as-2 and as-1 was then deleted. The petE promoter directs cell-specific but not full light-regulated expression in transpence tobacco plants. The promoter is used to construct chimeric promoters of the invention. The
                                                                                                                                                                                                                                                                                                                                        for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, meion, cucumber, pea, rape, beetroot, or sunflower, and monocotyledonous species, e.g. wheat,
                                                                                                                                                                                                                                                                                                                                                                                                    specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operably or functionally linked upstream of a CAAT box, TATA box and transcription initiation site. The chimeric promoters are used in expression vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgeric plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; Fage 72; 83pp; English.
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                                                                                                                                                                                                                                                                                  Sequence 220 BP; 73 A; 60 C;
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GenCore version 4.5 Copyright (n) 1943 - 2000 Compugen Lid

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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result	Score	Query Match	Query Match Length DB	80	ΠĐ	Description
1	146	10.8	394	10	BE319260	8404504N 03781834
N	146	10 8	306	10	RT264788	
ω	146	10 A	413	10	BE249329	
4	146	10 8	422		HF315615	
5	146	10.8	439	10	BE318254	
6	146	10.8	442	10	HF636746	-
7	146	10.8	449	10	BF323973	BE323973 NF011H07F
8	146	10.8	455	10	BE323979	_
9	146	10.8	464	10	BF520824	
10	146	10.8	474	10	HE316929	BK316929 NFQ53H061.
11	146	10.8	476	10	вк318158	
12	146		480	ìò	HF635626	HF635626 NF109F03D
13	146	10.8	482	10	BE633611	BP633611 NF058C04D
14	146	10 8	512	<u>1</u> 0	BE316878	~
15	146	10.8	516	10	BF632144	BF632144 NF033F10D
16	146	10.d	520	,	BF513450	REPORTED NEGRESOAD
17	146	10.8	523	ø	AW775723	AW775723 EST334788

AW776926	AW776926		635	10.8	146	45
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	33.7		627		146	<u>م</u> ن
	В127015	10	613	0	146	41
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	HE3172	.10	593	10.8	146	36
	HF.2.181	10	590	0.	146	35
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	BE3249	11	574	₽	146	نیا
	BE3191	10	572	10.8	146	دی ده
	HF63725	;=	555	Œ	145	
	) нг641618	1	565	С.	146	30
	нкз1813	10	565	10.8	146	29
	BE3241	11	on AD	0.	146	83
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	BG45528	15	546	10.8	146	20
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## ALTONMENTS

	Source	FEATURES	Sec	21d	Ins	<b>X</b> O:	Fau	T'c.	r.,	3G.E.	£1a	Cor	COMMENS On	JOURNAL Uni	Med	TITLE Exp	, G.			KEFERENCE 1	Med	Ros	Spe		ORGANISM Med	SOURCE bar	KEYWORDS EST.	VERSION BE	ACCESSION BET	NF(	DEFINITION NEG		BE319260	RESULT 1
/organismmenicago Lubratuid" /db_xref-mtaxon:3880" /rlone="NF016B04RT" /rlone_lib "Developing root" /tissue_type "root"		Location/Qualifiers	Seq primer: "CACACAGGAAACAGCTATGAC.	Plate: 016 row: B column: 04		Modicans Concept This in the homospinon, Matter 14645	C: 580 221 /380	Tel: 580 221 7317	2510 Sam Noble Parkway, Ardmitte, OF 73400, USA	The Samuel Roberts Noble Foundation	Plant Biology Division	Contact: Paiva NI.	Om Jul 14, 2000 this sequence version replaced grigity(06).	Unpublished (2000)	Medicago truncatula root library	Expressed Sequence Tags from the Samuel Roberts Noble Foundation	,G.D. and Paiva,N.L.	Gonzales, R.A., Bell, C.J., Flores, H.R., Inman, J.T., Weller, J.W., May	Son, B.S., Shin, HS., Lopez Meyer, M., Scott, A.D., Harris, A.R.,	l (bases 1 to 394)	Medicago.	Rosidae; eurosids I; Fabales; Fabaceae; Papilionoldeae; Tritolieae;	Spermatophyta: Magnoliophyta: eudicotyledons: core eudicots:	Bukaryóta; Viridiplantae; Streptophyta; Embryophyta; Tracleophyta;	Medicago truncatula	barrel medic.		BE319260.2 GI:11930467	BE319260	NF016H04RT 5', mRNA sequence.	NF016804RT1F1032 Developing root Medicago truncatula cDNA clone	BE319260 394 bp mrnA linear ESI 20 DEC 2000		

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CLORICA MELLEGUAPI. SE, E ENA SOSPHENCO.
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The Samuel Roberts Noble Foundation
2510 Sam Noble Parkway Ardmore, OK 7802, USA
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period. The points were tertilized twice weekly with 1/2 Hood ands soot from containing only ZOMM potassium presidente. ENA was prepared from above ground tissues."

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NFOLISCOLLFIELOOG beveloping leaf Medicado trancatula CONA clone-NEOLEGOLLF 57, mKNA sequence.
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The Samuel Roberts Noble Foundation
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Seq primer: FCACACAGGAAACAGCTATGAC
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Tel: 580-221-7391
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At Saue type "leat"

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/db xref "taxon:4880"
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The Samuel Roberts Noble Foundation
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on Jul 14, 2000 this sequence version replaced gi จำหนังชีวิ
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BE315615
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/tissue_type="leaf"
/dev_stage-"Pooled developmental"
/note-"Vecetor: Lambda Zap, Centains a mixture of very
young, developing, mature and senescing leaves."
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The Samuel Roberts Noble Foundation
2510 Sam Roble Parkway, Ardmore, OK 73402, USA
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The Samuel Koberts Noble Foundation
2510 Sam veble Parkway, Ardmere, oK J 402, USA
Tel: 580 221 7325
Email: mjharrison@noble.ord
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Tel: 612 625 1243
Eax: 651 649 5056
                                                                                                                                                                                                                                                                                                                                   Hoit, I.E. and Fraser, C.M. ESTs from leaves of Medicago truncatula after inoculation with Colletotrichum trifolii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Medicago truncatula
Eukaryota: Viridiflintae: Streptophyta, Embryophyta, Tracteophyta:
Spermatophyta; Magnoliophyta, eudicotyledons, core codicots;
Rosidae, curosids I, Fabales, Fabaceae, Papilionoideae; Trifolioao;
                                        University of Microsota name: M278256e
TIGR sequence name: MFPDA14TK
More information is available at. http://chrysic.tamu.edu/medicago
Seq primer: SKmod (CTA gAA CTA gtg gAT CC).
                                                                                                                                                                                                                                     Department of Plant Pathology University of Mirnesota
                                                                                                                                                                                                                                                                                     Unpublished (2000)
Contact: Deborah A. Samac
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/db_xref="taxon-3880"
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The Samuel Roberts Noble Foundation
2510 Sam Noble Parkway, Ardmore, CK 73402, USA
                       Torres-Jerez, I., Scott, A.D., Harris, A.R., Gonzales, K.A., Hell, C. Flores, H.R., Imman, J.T., Weller, J.W. and May, G.D. Expressed Sequence Tags from the Samuel Roberts Noble Foundation
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XhoI; cDNA was prepared from polyA- enriched RNA from
cotyledons and primary leaves harvested 5 and 8 days after
inoculation with Colleichtrichum trifolli. The cDNA was
directionally ligated into the Unit ZAP XR vector from
Strategiese and packaged using Cidapack [i] Cold packaging
extracts. Plasmids containing cDNA inserts were excised
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/lab_host="E. coli strain XLOLR"
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The Samuel Roberts Notice Foundation
2510 Sam Noble Parkwip Ardmere, eg.7 402, USA
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Tel: 580-221-7391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: gdmay@noble.org
Insert Length: 482 Std Error 0 |
Plate: 058 row: C COLUMN: 04
Seg primer: TCACACAGGAAACAGCTAFGAG
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Tel: 580 221 7391
Fax: 580 221 7380
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Uncry Match 10.8%: Score 14: DB 10: Best Local Similarity 100.0%: Pred. No. 1.96-42: O: Mismatches 0:
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Eusidae; eurosids I; Fabaies; Fabaceae; Papiliopoldeae; Trilolieae;
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/note "Vector: Lambda Zap, Fontalis a mixture of
young, developing, nature and senessing leaves."
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Contact: May Ch
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The Samuel Rober's Noble Foundation
2510 Sam Noble Parkwass Ardmores OK 74022 USA
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APPLICANT: Rice, bouglas A
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                                                                                                                                                                                    INFORMATION FOR SECTIONS:
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APPLICANT: Jian Ni and Jing Shan Hi
TITLE OF INVENTION: Human Georg, Sequences and
TITLE OF INVENTION: Express. I Products
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Best Local Similarity 100,c%; Prod. No. 12;
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INFORMATION FOR SECTIONO: 1:
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TOPOLOGY: Linear
MOLECULE TYPE: Boptide
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1446 ATGAATTGATGAAAGAGTT 1464
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CLASSIFICATION 10ATA: 435
PRI R APPLICATION NUMBER: 08/366,480
FILING DATE: DECOMBER: 08, 1994
CLASSIFICATION: 435
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LELEPHONE: (517) 847 4 100
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REGISTRATION NUMBER:
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                       Fest Local Similarity 100.0%;
            GENERAL INFORMATION:

APPLICANT: Sylvain Moincau, Shirley A
APPLICANT: Walker, Ebenezer R. Vedamuthu,
APPLICANT: and Peter A. Vandenbergh
TITLE OF INVENTION: Isolated DNA Encoding
TITLE OF INVENTION: Enzyme For Phage
TITLE OF INVENTION: Resistance
                                                                                                                                                                                                                                                                             1446 ATGAATTGATGAAAGAGTT 1464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 1:
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TELEPHONE: (517) 347-4103
TELEPHS: (517) 347-4103
TELEPA: No. 5925388e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: MS-DOS
SOFTWARE: WOTOPERFOCE 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: TS/08/820,980
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MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb
MEDIUM TYPE: storage
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ADDRESSEE Tan C McLand
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TITLE OF INVENTION:
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APPLICANT: Madker, Ebenezer R. Vedanuthu,
APPLICANT: and Peter A. Vandenbergh
TITLE OF INVENTION: Isolated DNA Encoding
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/424,6
FILING DATE: April 19, 1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 20,931
REFERENCE/DOCKET NUMBER: Quest 4.1-156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE:
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Resistance
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100.0%; Pred. No. 17;
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US-08-913-159-1
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; Patent No. 6300109
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                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                              APPLICATION NUMBER: DK 91
FILING DATE: 17-FEB-1995
INFORMATION FOR SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1446 ATGAATTGATGAAAGAGTT 1464
                                                                                                                  REFERENCE/DOCKET NUMBER: OUTELECOMMUNICATION INFORMATION:
TELEPHONE: (517) 347-4100
TELEPAX: (517) 447-4103
                     SEQUENCE CHARACTERISTICS:
LENGTH: 3695 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30 (EPG) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Plasmid-derived type II
TITLE OF INVENTION: restriction modification systems from Lactococcus Lactis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:

CENGTH: 2987 base pairs

TYPE: Nucleic Acid

STRANDEDNESS: Double
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ATTORNEY/AGENT INFORMATION:
NAME: Ian C. McLeod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SCFTWARE. Wordperfect 5.1 CURRENT APPLICATION DATA:
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ADDRESSEE: Ian C. McLood
                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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APPLICATION NUMBER: 08/424;
FILING DATE: APPTI 19, 1995
CLASSIFICATION: 530
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CITY: Okemos
STATE: Michigan
                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEX: No. 59726734
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OPERATING SYSTEM: MS-DOS
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                                                                                                                                                                  FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/826,439 FILING DATE:
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    STRANDEDNESS:
                                                                                                                                                                                      APPLICATION NUMBER: US/08/913,159
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Hest Local Similarity 100_0%; Pred. No. 17;
                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
             APPLICATION NUMBER:
                                                                                  SOFTWARE: Patent in Release #1.0. Version #i.40 (390) CURRENT APPLICATION DAFA.
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Mot Bankar annibanar
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                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                           FITLE OF INVENTION: restriction modification systems from Lactococcus Lactis
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                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC computibles
OPERATING SYSTEM: PC 1905/METERS
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offHer information. /cools.start information. /cools.start information. /cools.start information. /cools.start information. /cools.good.
FILLING DATE:
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                                                                     APPLICATION NUMBER:
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/labet millaAfR
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LILLE OF INVENTION: restriction modification systems from Lactococcus Lactos
NUMBER OF SEQUENCES: 14
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MEDIUM TYPE: Floppy disk
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IDENTIFICATION METHOD:
OTHER INFORMATION: /cod
OTHER INFORMATION: /cod
OTHER INFORMATION: /cyd
OTHER INFORMATION: /cyd
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                                                                                                                                                                                                      FEATURE.
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                                                                                                                                                                                                                                                                                                                SEIGINAL SOURCE:
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OTHER INFERMATION: /codom_start: [649]
                                                                                                                                                                                                                                           SIRAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERALING SYSTEM. PC DESIME 1835
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                                                                             IDENTIFICATION METHOD: experimental
                                                                                                                         LOCATION:
                                                                                                                                                          NAME/KEY: CDS
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STRANDEDNESS: double
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190,0%; Pred. No. 17;
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/product="LIAAL MAIO ademine methyluse H"
/cyldence_EXPERIMENTAL
/qenc=""BRE"
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OTHER INFORMATION:
OTHER INFORMATION:
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US-08-913-159-5
Дb
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NEORMATION FOR SEQ ID NO: 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO) CURRENT APPLICATION DATA:
2118 ATGAATTGATGAAAGAGTI 2136
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OTHER INFORMATION: /pr
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OTHER INFORMATION: /ge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION.
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                               515 atgaattgatgaaagagtt 533
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OTHER INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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LOCATION: 2412.
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100 0%; Pred No
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                                                                                      100 0%;
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                                                                                                                                                                       /standard_name- "Gene coding for LlaAI restriction endonuclease" /label= r-llaAI
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/gene- "ORF"
/number- 2
/s andard_name- "Gene coding for M.LlaNIB"
/label- m-llaNIB
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US-09-128-155-16/c
; Sequence 16, Application "S/09128155
; Patent No. 6117654
                                                                                                                                                                                                      ; NAME/KEY: misc_feature
; LOCATION: (1)...(176373)
; OTHER INFORMATION: n - A,T,C or G
US-09-128-155-17
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: Sequence 17, Application US/09128155
: Patent No. 6117654
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; LUCATION: (1)...(15231)
; OTHER INFORMATION: n = A,T,C or
US-09-128-155-16
Db 108094 aagaacaatgctggttttc 108112
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                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 17
                                                                                             Matches
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Best Local Similarity 100.0%;
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TITLE OF INVENTION: NOVEL MOLECULES OF TANGO-77 RELATED PROTEIN FAMILY
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: 0444,705,001
CTREET APPLICATION NUMBER: 08/09/128,155
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EARLIER FILING DATE. 1997 08-04
NUMBER OF SEQ ID NOS: 18
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TITLE OF INVENTION: NOVEL MOLECULES OF TANGO-77 RELATED PROTEIN FAMILY
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: 69404/652001
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EARLIEF FILING DATE: 1997-98-04
NUMBER OF SEQ ID NOS: 18
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100.0%; Pred. No. 14;
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100.0%; Pred. No. 13;
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RESULT 15
US OB 78 588 15
Sequence 15. Application USZOB COURT
Datent no. 5608 148
GENERAL INFORMATION:
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: LOCATION: (1)...(922)
(6: 09-47/-648-4
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Best Local Similarity 19-18: Pred. No. 50:
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APPLICANT: Martino Cart, Susai
FILLE OF LOVENTION: Seed Frederied Promoters
FILE REFERENCE: 0469
CHRRENT APPLICATION NUMBER: US 509/447,648
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INFORMATION FOR SEQ ID NO: 15
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NAME SCAY, NICHOLAS J.
NAME SCAY, NICHOLAS J.
REFERENCE/DOFKET NUMBER: 11
TELECOMMUNICATION INTERMATION:
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## ALIGNMENTS

DEFINITION ACCESSION VERSION KEYWORDS RESULT AX105268 LOCUS ORIGIN BASE COUNT FEATURES REFERENCE SOURCE TITLE JOURNAL ORGANISM AUTHORS Vezina,L.P. and D'Aoust,M.A.

Promoter for regulating expression of foreign dense patent; WO 0125455-A 2 12-APR-2001;

WEST-RESERVED TO CASE AX105268 Sequence 2 from Patent W00125455 AX105268 AX105268.1 G1:13921418 artificial sequence. 1 (bases 1 to 971) synthetic construct Medicago Inc. synthetic construct 9 () F /organism="synthetic construct"
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174. .205
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Local Similarity 100.0%; Pred. No. 3.1e-16;
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Spermatophyta; Magnoliophyta; eudicotyledons; core cud cots;
Rosidae, eurosids I, Fabales, Fabaccae, Papilionoideae, Victeae,
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                                                                                                            /product-"mature plastocyanin (AA 1-99)"
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813. .1319
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59. .88
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25. .62
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/db_xref="taxon:3888"
/clone="lambda-GPC3"
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134, 171
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130. .159
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21. .50
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aslkdegyatvataasavlasnalavevllgasdgglafvpsslevsagetivfknna
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                                                                                                                                                                                                                                                                                 /protein_id-"CAA34212 1"
/db_xref-"G1:20846"
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                                                                                                                                                                                                                                                                                                                                      /note-"precursor peptide (AA -69 to 99)"
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Rance, I., Theisen, M. and Gruber, V.
Patent: FP 2701358-A 6 29-5FP-2000;
                                                                                                                                                                                                                                                                                                                                                                              Sequence 6 from Patent FR2791358
Ax036600
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                                                                                                                                                                                                                                                                                                                       synthetic construct
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Patent: PR 2791358-A 13 29-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           artificial sequence.
1 (bases 1 to 68)
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                                                                                                                    /note-"The promoter MPrII09 differs from MPrII09 by a deletion of 33 bp in the 5' UTR, II bp upstream of the point +1
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                                                                                                      promoter MPr1109"
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/db_xref="taxon:32630"
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100.0%; Pred. No. 2.9e-05
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Local Similarity (100 0%) Pred. No. 0.000/6;
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                                                   41; Conservative
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AXU66601
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1 (bases 1 to 280)
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AX036598
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promoter Mpr109H"

a "B = 21 g & 61 t
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                                                                                                                                                           promoter MPF1110"
4 24 · 42 u
                                                                                                                                                                                        CaMV promoter
                                                                                                                                                                                                       position 99 bp of Meri098 an element of 18 bp containing a 55 bex and by fusing a Sequence of 44 bp of the KNA 85S
                                                                                                                                                                                                                                        /note "The promoter MircillO was created by inserting at
                                                                                                                                                                                                                                                                           /organism "synthetic construct"
/h-xcal "tasan-quam"
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Leal Similarity 198, %; Bred. No. 10,08070;
Hes 31; Conservative 0; Mismatches 0; Indeed
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                                                                                                                                                                                     Rance, I., Theisen, M. and Gruber, V. Patent: FR 2791358 A 8 29 SEP 2000: MERISIEM (HERAPEULL'S (FR)
                                                                                                                                                                                                                                                                        synthetic construct
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AXO36603
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Sequence 8 from Patent FE2791358.
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            promoter Merilian
                                              sequence of 28~\mathrm{hp} of the promoter peth, stretching true position, 682~\mathrm{to} position. Six by modified by adjunction
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a 74 c 4% a
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                                  And it to the
                                                                                 /mote "The promoter MPT1153 was obtained by tusing
                                                                                                        /db vret "taxon.326 d)"
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## ALIGNMENTS

RESULT AAS02127 ID AAS0 Novel promoter for regulating expression of foreign genes organisms, more specifically in a leaf specific manner in 04 - OCT - 1999; Promoter #2 for regulating expression of foreign genes AASU2127 standard; DNA; 971 BP WPI; 2001 266316/27 Vezina L. 02-OCT-2000; 2000WO-CA01144 12 - APR - 2001 WUZ00125455-A2 Fromoter, transpesse plant, munocotyledom; dicotyledom; dymnosperm; ds 18-JUn-2001 AAS02127: (MEDI-) MEDICAGO INC. \_ D'Aoust M; (first entry) 990S-0157129

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              Sucry Match
Best Local Similarity
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                     Chimeric expression promoter to producing dreatyledomous and monocotyledomous transferic plants comprises a nucleic acid septence derived from a promoter of the pea plastocyanin dene
                                                                                                                                              site. The chimeric promotors are used in expression vectors for
producing transgence plants, such as dicotyleborous species,
e.g. potato, tabarco, certon. Lettuce, tamato, melon, encumber, pea-
                                                                                                                                                                                                describes a chimeric expression promoter comprising a promoter of the pea plastocyanin deno, or comprising a 6-box operably or functionally linked upstream of a CAAL box. TATA box and transcription initiation
                                                                                                                                                                                                                                                                                                      The present sequence represents the chimeric promoter Mpr1096. The promoter is derived from the petE promoter from pea plastocyanin gene, by a deletion in 5° of the repeated invert sequences, of the as I like and enhancer like elements beine by two Sphl frauments of 403 bp. The
                                                                                  Sequence 326 BP: 111 A: 31 3
                                                                                                                    barley, oat, rice, or cons.
                                                                                                                                    rape, beetroot, or sunfilewer, and monocotyledomous species, e.g. wheat,
                                                                                                                                                                                                                                                      construct chimeric promoters of the invention. The specification
                                                                                                                                                                                                                                                                    expression in transpenie tobasco plants. The promoter is used to
                                                                                                                                                                                                                                                                                                                                                                                        Claim 2: Page 67: 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI: 2000 587667755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rance L. Gruber V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MERL ) MERLSTEM THERAPPOIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22 MVB [1884]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0000 JHS 87
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 petE promoter: chimeric promoter: transferic plant; MPr1096,
plastocyanic dene promoter: s;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleatide sequence of the promoter MPr1396
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA96459;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA95459 Standard; DNA; (25)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      961 qaqaqaaaatq 971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         721 gagagaaaaata 731
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Score 57: DB 21; Pred. No. 1.6e-14; 0: Mismatches =:

Length 523

1,985

MESULI AAA95457

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AAA96457 standard: DNA; 834

ΔΔΔ36457

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expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a 6 box operably of functionally linked distream of a PAN box. TAIA hox and transcription initiation site, the chameric panches are used in expression vectors for producing transaction plants, such as disortyledonous species, e.g. potato, tobacco, colton, lettuce, tomato.
                                                                                                                                                                                                                                 promoter is derived from the petE promoter from pea plastocyanin are
by a deletion in 5′ of the repeated invert sequences as well as the
as 1 like box berne on a SphI fragment of 212 bp. The petE promoter
directs cell-specific but not full light regulated expression in
Sequence 423 BP: 230 A: 128 C: 82 C: 183 I: 0 other
                                                              melon, cucumber, pea, rape, beetroot, or sunflower, and monocotyledonous
                                                                                                                                                                                              transferic tobacco plants, the promoter is used to construct enumeric promoters of the invention. The specification describes a chimeric
                                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 65; 83pps English
                                                                                                                                                                                                                                                                                                                                                                                                                                       Chineric expression promoter for producing directyledonous and monagentyledonous transquaric plants comprises a nucleic acid sequence
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                                         species, e.g. wheat, barley, oat, rice, or corn
                                                                                                                                                                                                                                                                                                                               The present sequence represents the chimeric promoter Mp.1897.
                                                                                                                                                                                                                                                                                                                                                                                                                     derived from a promoter of the pea plasforyanin dene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPT: 2000 587662/55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MEEL ) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22 MAR 1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 MAR 2000; 2000WO IBOUT!/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28 SEP 2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fisum Sativum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleotide sequence of the promoter MPr1097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UN FEB 2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA96458;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         petE promoter; chimeric promoter, transdemic plant: Micle97;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ٦
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           33 SECTION 836.65
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Best Local
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                                                                                                                                                                                                                                         6468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a 0 kew specially or functionally linked upstream of a CAAF box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous species, a.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, beetroof, or sunflower, and monocetyledonous species.
                                                                                                                                 Nucleat ide
                                                                                                                                                            08-FEH-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 834 BP; 289 A; 175 C; 133 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               species, e.g. wheat, barley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents the petE promoter from the pea plastocyanin gene. The promoter directs cell-specific but not full light-regulated expression in transgenic tobacco plants. The promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 65; 83pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        plastocyanin gene promoter; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleotide sequence of the petE promoter from plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-FEB-2001 (first entry)
28-SEP-2000
                            W0200056906-A1
                                                       Synthetic
                                                                                     plastocyanin
                                                                                                 petE promoter; chimeric promoter; transgenic plant; MPr1108.
                                                                                                                                                                                           AAA96468;
                                                                                                                                                                                                                        AAA96468 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      derived from a promoter of the pea plastocyanin gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        petE promoter, chimeric promoter; transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 is used to construct chimeric promoters of the invention
                                                                                                                                                                                                                                                                                             618 agataacccactttaagcccacqcactctgtggcacatctacattatclaaatcaca 674
                                                                                                                                                                                                                                                                                                                         759 agataacccactttaagcccacgcactctgtggcacatctacattatctaaatcaca 815
                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                              Gruber V,
                                                                                                                               sequence of the directional building block S3.
                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                   promoter; PCk primer:
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                                                                                                                                                                                                                                                                                                                                                                       Score 57;
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                                                                                                                                                                                                                                                                                                                                                                                                                                237 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                       DR 21;
                                                                                                                                                                                                                                                                                                                                                                                     Length 834:
                                                                                                                                                                                                                                                                                                                                                         Tidelig
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Best Local Similarity
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Chimeric expression promoter for producing dicotyledonous as measuryledonous transgenic plants comprises a nucleic acid derived from a promoter of the pea plastocyania gene.
                                                                                                                                                                                                                                                                                                                                                               Nucleotide sequence of the promoter MPr1109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plants, such as dicutyledonous species, e.g. potato, tobacco, lettuce, tomato, meion, eucumber, pea, rape, bectroot, or sun and monostotyledonous species, e.g. wheat, bariey, oat, rice, o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprising a G box operably or functionally linked upstream of a CAAT box, TATA box and transcription initiation site. The chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       promoter comprising a petE promoter of the rea plastocyanin gene, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 24; Page 73; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
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                                                              WPI; 2000-587667/55
                                                                                         Rance I,
                                                                                                                                                     22-MAR-1999;
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                                                                                                                                                                                                                                                                        Pisum sativum
                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                    plastocyanin
                                                                                                                                                                                                                                                                                                                                                                                               UB-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                             AAA96462;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA96462 standard; DNA; 250 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             promoters are used in expression vectors for producing transquaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       which is used to construct chimeric promoters of the invention in PCR reactions. The specification describes a chimeric expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rance I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                   petE promoter; chimeric promoter; transgenic plant; MPr1109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              782 dadtotytyyddadatotadattatotadatodda 815
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present sequence represents the directional building block
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MERISTEM THERAPEUTICS
                                                                                         Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ilarity 100.0%;
Conservative (
                                                                                                                                                                                  2000WO-1800317
                                                                                                                                                                                                                                                                                                                    gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99FR-0003635
                                                                                                                                                    99FR-0003635
                                                                                                                                                                                                                                                                                                                    promoter;
                                                                                         Theisen M.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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9e-05;
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Matches
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                 minimal reference promoter of the petE promoter. The petE promoter directs cell specific but not full light required expression in
                                                                           the present sequence represents the chimeric promoter Mp.1698. The promoter is derived trem the petE premeter trem per plantacyanin gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            producter is derived from the petE producter from per plastoryania dense, by fusing the petE as 1 lake and most enhancer lake elements to the producter Mut1098 (compressing FALA and CAAI boxes of petE), and then deleting 4 bp from the 5 U.R. The petE producter directs cell specific deleting 4 bp from the 5 U.R. The petE producter directs cell specific
transgenic tobacco plants. The promoter
                                                            and contains only the TAIA and CAAT bexes, and corresponds to a
                                                                                                                                          Claim 2: Page 62, Bipper Shalinda
                                                                                                                                                                                   derived from a promoter of the pea plasforyanin dene
                                                                                                                                                                                                         monocotyledonous transaedic frants comprises a nucleic acid sequence
                                                                                                                                                                                                                            Challetic expression productor to producing disotyledonous and
                                                                                                                                                                                                                                                                     WP1; 2000 587552/35
                                                                                                                                                                                                                                                                                                               Rance I, Grober V.
                                                                                                                                                                                                                                                                                                                                                   (MERT ) MERISTEM THERAPLETE'
                                                                                                                                                                                                                                                                                                                                                                                            22 MAR 1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     plastocyanin dene promoter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 250 BP; 81 A; (20); (1 G; 71 1, 0 other)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         species, e.g. wheat, barley, sat, rice, or corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          dieotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, per, rape, leetroot, or suntlower, and monocotyledonous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression vectors for producing transgenic plants, such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The promoter is used to construct chimeric promoters of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      but not full light requiated expression in transgenie tobacco plants
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2: Page 69 Bipp: Eng. sh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        petE promoter: chimerre promoter: transgenic plant; MPr1897;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               specification describes a chimeric expression promoter compaising
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  as used to construct chameric
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Best Local 1
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                                                                                       transpenie tobacco plants. The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric expression promoter competished a parameter of the party-train of a "AAI box and transcription institution site. The chimeric promoters are constructed as a consequence of the party-train of a "AAI box and transcription institution site. The chimeric promoters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                promotets of the invention. The specification describes a chameric expression promoter comprising a posteryach decorporation promoter at the peal posteryach decorporation is box operably or functionally linked distinct or a CAN box. TATA box and transcription initiation site, the chameric promoters are used in expression vectors for producing transactic plants, such as disotyledonous species, e.g. potato, tobacco, estical estimate, tomato.
                                                                                                                                                                                                                                                    promoter is derived from the petE promoter from pea plastocyanin dene, by fusing the petE as 1 like and nos enhancer. Tike elements to the promoter MYT1098 (comprise Tid Tala and CAAT baxes of petE; and fixed inserting a 6 box and fissing a frammental des NNA 22 PMV. The patE promoter directs cell specific but not full light regulated expression in
are keed in expression sectors for producing transquaic plants, such as directyledonous species, e.g. potato, tobacco, cotton, between tomato, melon, cucumber, pea, rape, bestroot, or sunflower, and homocotyledonous
                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents the chimeric promoter Mprillo. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Maim 2; Page 70; 83pp. Egglish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric expression promoter for producing dicotyledomous and monocotyledomous transuenic plants comprises a nucleic acid sederized from a promoter of the pea plastocyanin dene-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-MAR 1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W0200056906-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide sequence of the promoter Mirelilia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAAS6463 standard; (NA; 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Séquence 207 BP; 67 A; 58 C; 21 G; 61 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    dicotyledonous species, e.g. potato, tobacco, cutton, cutture, tomato, medon, cucumber, pea, rape, bectroot, or suntrower, and namonetyledonous species, e.g. wheat, barbay, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MERI ) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAR 2000; 2000Wo [B004]7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA96455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         petE promoter: chimerie promoter: transdence plant: METILIO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    784 etetatogeacaletacaltaletaaaleaca 815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lf - hitataa aa atitacattatotaaatcaca 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000 587667/55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gruber V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dene bromoter:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39FR 0003545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      [160]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 4 4
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Prod. No. 0.00058;
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species, e.g. wheat, barb $\gamma_i$  oat, nice, o

os O

Sequence 280 HP; 88 A; 73 C; 42 G; 77 T; 0 other

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AAAS6465
ID AAAS
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                  Matches
                                                                                                                                                                                   promoter of the pea plastocyanin gene, or comprising a G box operably or functionally linked upstream of a GART box, TATA box and transcription initiation site. The crimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledorous species, e.g. potato, tobacco, cotton, lettuce, tomato, melor, cucumter, pea,
                                                                                                                                                                                                                                                                                           inserting a G box and fusing a fragment comprising a duplication of the element as2 and as1. The pott promoter directs cell-specific but not full light-regulated expression in transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The
                                                                                                                                                                                                                                                                                                                                                               The present sequence represents the chimeric promoter Mpilllll. The promoter is derived from the petE promoter from ped plastocyanin gence by fusing the petE as-1 like and nos enhancer like elements to the promoter MPr1098 (comprising TATA and CAAT boxes of petE), and then
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2; Page 71; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-587667/55
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                                                                                                                                                                                                                                                                           specification describes a chimeric expression promoter comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rance I, Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-MAR-2000; 2000WO-IB00317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-SEP-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plastocyanin gene promoter; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           petE promoter, chimeric promoter, transgenic plant; MPr1111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleotide sequence of the promoter MPr1111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA96465 standard; DNA; 296 BP.
                                                                                                                      Sequence 296 BP, 94 A, 74 C, 45 G; 83 T; 0 other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MERI-) MERISTEM THERAPHUTICS.
                                                                                                                                                      barley, oat, rice, or corn
782 nacticitytygnacatni snattatotadato 912
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        68 cactctgtggcacatctacattatctaaatc 98
                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            beetroot, or sunflower, and monocotyledonous species, e.g. wheat,
                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                  Conservative
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                                                 100.0%;
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                                  <u>_</u>
                                                 Score 31;
Pred. No.
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Pred. No.
                                Mismatches
                                                                   DB 21; Length 296;
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0.0015;
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                                Gaps
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   Query Match
Best Local :
                                                                                                                                                                              describes a chimeric expression promoter comprising a promoter of the pea plastucyanin gene, or comprising a 6 kox operably or functionally linked upstroam of a CAAT box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing
                                                                                                                                                                                                                                                                                         The present sequence represents the chimeric promoter Mpr1153. The promoter is derived from the petE promoter from pea plastocyanin generated the petE promoter directs cell-specific but not full light-regulated
                                                                                                                                                                                                                                                                                                                                                                                                  Chimetic expression promoter for producing disobjectors and monocotyledonous transgenic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
                                                                      Sequence 303 BP; 101 A; 78 C: 41 G; 83 T; 0 other;
                                                                                                             or corn.
                                                                                                                                             cotton,
                                                                                                                                                                                                                                                                        expression in transgenie tobacco plants. The promoter is used to
                                                                                                                                                                                                                                                                                                                                                                 Claim 2; Page 70; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000 587667,755
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-FEB-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA96464 standard;
                                                                                                                                                              transgenic plants, such as dicotyledonous species, e.g. potato, tobacco
                                                                                                                                                                                                                                                        construct chimeric promoters of the invention. The specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rance I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         plastocyanin gene promoter:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        petE promoter; chimeric promoter; transgenic plant; MPr1153;
y Match 3.2%; Score 31; DB 21; Local Similarity 100.0%; Pred. No. 0.3015;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                84 cactetgtggcacatetacattatetaaate 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lettuce, tomato, melon, cucumber, pea, rape, beetroot, or er, and morecotyledonous species, e.a. wheat, barbe, wat, tire,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99FR-0003635
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                 Length 303;
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RESULT 13
AAA96470
ID AAA96470 standard; DNA: 72 BP.
XX
AC AAA96470;
XX
US FEB-2001 (first cetry)
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Matches

31,

Conservative

Ö

Mismatches

0

Indels

0

Gaps

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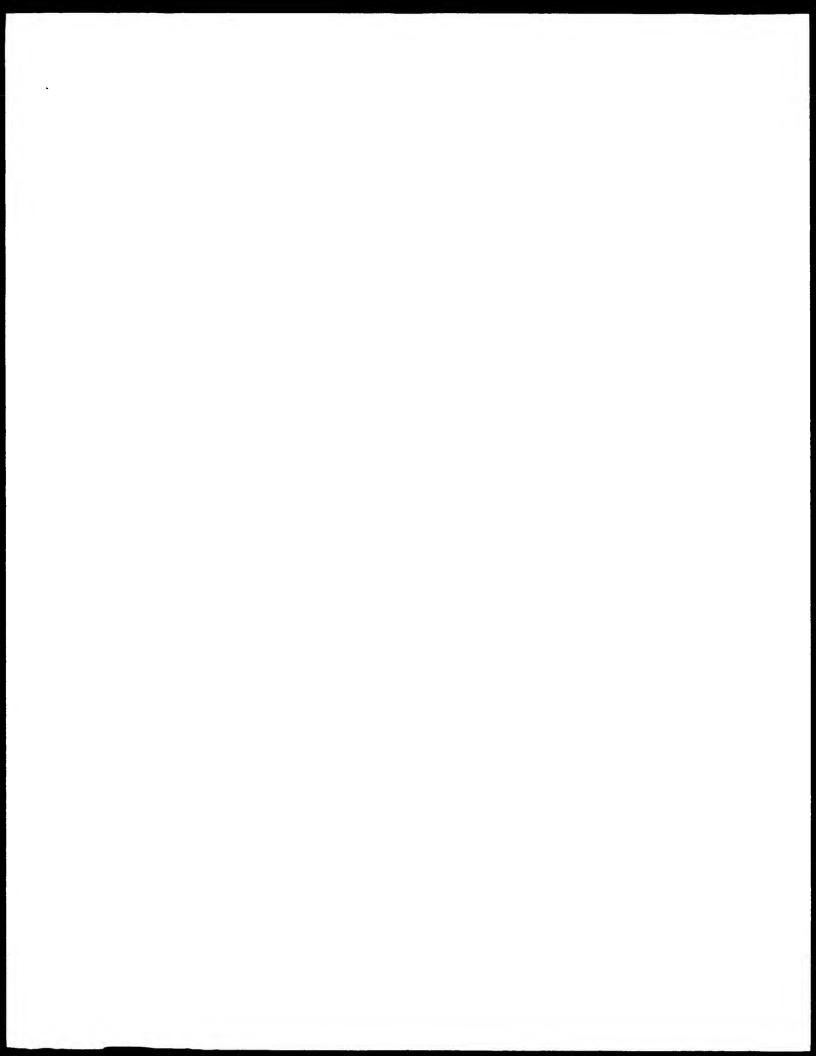
782 cactetytygeacatetacattatetaaate 812

91 cactotytyycacatotacattatotaaato 121

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\begin{array}{c} \mathbb{D} \times \mathbb{T} \times \mathbb{T} \times \mathbb{T} \times \mathbb{T} \times \mathbb{C} \times \mathbb{X} \times \mathbb{X} \times \mathbb{T} \times \mathbb{T} \times \mathbb{A} \times \mathbb{A} \times \mathbb{C} \times \mathbb{
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT / 7033
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OF HIN TYP!
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Best Local Similarity 100.7%; Pred. No. 0.07 c
Matches 27; Conservation 6; Mismatches 7; Indebs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the GUS reporter gene in tobacco where the GUS gene was under the control of the PalE promoter and 45% CAMV promoter. The enhancer was also used joined to the 340 to +1 region of the patatin promoter 1525 which was joined to the GUS reporter and transformed into potato using A.tumetaciens. Expression of the gene promoter of one of more genes of a plant increases in green of non-requirissnes, especially roots, tubers, seeds, stems, flowers or leaves, this enhances may operate both in normal and reverse directions and can be attached either to the promoter or terminator of the gene to be expressed.
WPI: 2000 587667/55
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BE532144 NE033E10D	BE315878 NEO57H08L	190M8504M 62691818	BC315615 NF024A08L	BI318452 NF038G07L	BE318195 NF036A09L	BE249329 NF013G01L	BE249398 NF013D12L	BL317478 NF070C071.	BE316779 NF067E07L	AW683569 NF016A041.	BE149600 NF023D09L	BF519460 BST456923	BF633855 NF065A06D	BF611829 NF014E071	dlübiloan kööktein	AL368181 MtHA22F11	BF636774 NF002E09L	BE317588 NF051H08L	AW777061 M110357e	AW127435 M310622 D	AK777901 M110695c	BE317561 NF051E04L	BE317607 NF054B11L	EI271170 NF051D03F	BI308417 EST529827	PENCHANCE FORMANCESA	BC586489 FST488257

# ACCESSION VERSION KEYWORDS SOURCE ORGANISM RESULT 1 BF635678/c LOCUS DEFINITION HF635678 375 bp mPNA linear ESI 19-DEC 2000 NF109F09D1F1677 DYOUGHT Medicago truncatula cDNA close NF109F09D1 5', mRNA sequence. BF635678 BF635678.1 GI:11899836 EST.

ALIGNMENTS

ORGANISM	ORGANISM Medicago truncatula
	Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
	Spermalophyta; Magnoliophyta; eudicotyledons; core eudicots;
	Rosidae: eurosids I: Fahales; Fahaceae; Papillionoideae; Tritolliege;
	Medicago.
FIFE KENCE	1 (bases 1 to 3/b)
AUTHORS	Torrer-Jerez, L., Spott, A.D., Barris, A.R., Contales, R.A., Bell, C.J.,
	Flores, H.R., Inman, J.F., Weller, J.W. and May, G.D.
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Plant Biology Division	TITLE TOURNAL COMMENT
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	source	FEATURES										COMMENT	TANALIOT
/organism="Medicago truncatula" /db_xrot="taxon:3880" /clone=lib="Drought" /tloss=t_tpe="Plantlets" /tlss=t_tpe="Plantlets" /det_stay="Proied timepoints" /tot="Vorton tarked time for the days-post-watering plantlets hardested in a scrips of days-post-watering	1376	Location, Qualifiers	Seq primer: TCACACAGAAGAACAGCTATGAC.	Plate, 109 row: Floodomn: 09	Hisert Length: 376 Std Error: 0.00	Email: gdmay/anoble.org	Tuk: 580 001 7380	Tel: 580 221 7391	2510 Sam Noble Pankway, Ardmore, OK 73401, HSA	The Samuel Roberts Nobic Foundation	Plant Biology Division	Contact: May GD	Unpublished (2000)

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Flores, H.E., Immaril, , Welfer, J.W. and May, G.B.
Expressed September Lies from the Samme. Roberts Noble Foundation
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NEOSOCIOFILES, mena sequence.
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The Samuel Roberts Noble Foundation
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147 c. 119 g. 174 t. Fothers.
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very young, developing, fully opened flowers and flowers
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                                                                  Medicago truncatala
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BC.X. Z., Chadle, C., Shadi, A.G., Barris, A.E., Sonrales, E.A., 10, C.J., Flores, H.R., Inman, J.T., Weller, J.W., May S.D., and Dixon
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2510 Sam Nober Forkway, Addmore, eK 241-.
Tel: 586-221 7460
Fax: 580-221 7460
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/db_xref "taxon:3880"
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                                                                                                               2510 Sam Noble Parkway, Ardmore, OK
Tel: 580 221 7391
Fax: 580 221 7380
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                   Email: qdmay@noble orq
Insert Length: 369 St.d Error: 0 f
Plate: 022 row: H rollin: 25
Seq primer: TCACACAGGAAACAGCTATGAC
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The Samuel Roberts Noble Foundation
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/clone_fib="Developing leaf"
/tissue_type="leaf"
/dev_staye="Pooled developmental"
/note="Vector: Lambda Zap, Contains a mixture of very
young, developing, mature and semesting leaves "
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:9:
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Insert Length: 378 Std Error: 0.1
Plate: 002 row G column: 02
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Spermalophyla, Magsoliophyla, endicolyledoka, core endicols;
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                                                                                                                                                                                                                                                                                                                                                                 2510 Sam Noble Parkway, Ardmore, OK 33402, USA
Tel: 580 221 7391
Fax: 580 221 7380
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Expressed Sequence Tags from the Samuel Roberts Noble Foundation

Medicago truncatula flower library
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/close_ib "Developing flower"
/tissue_type "beveloping flowers"
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/dev_stage "beveloping, fully-opened flowers and flowers
in early transition into pods."
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2510 Sam Noble Parkwa. Ardmote, 58-2-602. USA
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Seq primer: FCATACAGAAACAGTATGAT.
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,A.J., Flores,H.R., Imman,J.L., Weiler, L.W., Mag. 1.0. and Joseph
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Tel: 580-221-7391
Fax: 580-221-7380
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Email: mjharrison@noble.org
Insert Length: 647 Std Error: 0 00
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Seq primer: TCACACACAGGAAACAGCTATGAC
                                                                                                    The Samuel Roberts Noble Foundation 2510 Sam Noble Parkway, Ardmore, OK 73402, USA Tel: 580 221 7325 Fax: 580 221 7380
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Spermatophyta, Magneliophyta, eudicotyledons, Norweudicous,
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/clone_type-"Devalping leat"
/tissuc_type="leaf"
/dev_stare "Pooled developmental"
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Eukaryota, Viridiplantae, Streptophyla, Embryophyla, Tracheophyla;
Spermatophyta, Magnoliophyla, eudicotyledons, core eudicots;
Spermatophyta, Magnoliophyla, eudicotyledons, core eudicots;
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NF014009LF1F1059 Developing leaf Medicago truncatula cDNA clone
NF014009LF 5', mRNA sequence.
                                                                                                                                                                                                                                                                            Email: qdmay@noble.orq
Medicaqo aerome Trittative accession
Insert Length: 645 Std Error: 0.00
Plate: 014 row: C column: 09
Seg primer: TCACACAGGAAACAGCTATGAC.
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Similarity 100.0%; Pred. No. 3.9;
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a 148 c 128 g 182 t l others
/dev_stage="Pooled developmental"
/note-"Vector: Lambda Zap, Contains a mixture of
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                                                                                                     /clone="NF014C091.F"
/clone_lib="Developing leaf"
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/hoto "Vector, (ambda Sap: At the trifoliate stage, M.
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/tissue_type="leaf"
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Torrestlerez, L. Scott, A.D., Barris, A.R., Ponzalet, R.A., Bellott, J. Flores, H.R., Tuman, J.E., Wellet, J.W. and May, G.D. Expressed Sequence field from the Manual Roberts Noble boundation
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Department of Plant Pathology
University of Minnesota
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,H., Ellis,L., Town,C.D., Bowman,C.L., Chree, M.R., Hansen,I.S.
Hott,C.F. and Fraser,C.M.
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/oulfiyar "acondigo Al/"
/db_xret-"iaxon 6860"
/clone "pDSIL 17817"
/clone_lib "DSIL"
/rissue_rype-"leaves intected with Scilestrictum
frifoli]
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RESULT 15
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T TLE
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Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM
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On Apr 14, 2000 this sequence version replaced gi.7558361
Contact: May Gh
                                                                                                                                                                                                                                      Texas A&M University
Department of Plant Pathology and Microbiology, Rm 120 L.F.
Peterson Bidg, College Station, TX 77843-2132, USA
Tel: 409 845 874;
Fax: 409 862 4790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A0841643 468 bp DNA linear GSS 27-SEP-1999 T137356b shotgun sub-library of BAC clone 72F02 Medicago truncatula genomic clone 72-F-02-084, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: gdmay@noble.org
Insert Length: 713 - Std Error: 0.00
Plate: 016 - row: 3 - orlumn: 10
Seq primer: TCACACAGGAAACAGCTATGAC.
                                                                                                                  Seq primer: pUC-C
Class: BAC subclone.
                                                                                                                                                     BAC survey sequencing of Medicago truncatula (1999c) Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                       The Crop Biotechnology Center
                                                                                                                                                                                                                                                                                                                                                                Contact: Cook DR
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/note="vector: Lambda Pap: Centains a mixture of
young, developing, matter and senescing leaves."
a 85 c 75 g 101 t
              /organism-"Medicago truncatula"
/cultivar-"genotype A17"
/db_xre!="taxon:3880"
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/tissue_type-"leaf"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:3880"
/clone="NF016G10LF"
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/clone- '72-F-02-084"
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/clone_lib-"shotgun sub-library of BAC close 72F02"
/note-"Vector: pUC18; BAC survey sequences were obtained from sheared BAC DAA subcloned into the Smal site of pUC18. The template DNA for sequencing was obtained by PCk using universal primers. Sequencing reactions were primed from the PUC-C primer site (CASSAAACASCTATGACCATGATTACGA) in the PUC18 polylinker; to the PUC18 puch the PU
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Query Match 2.9%; Score 28; DB 12; Length 468; Best Local Similarity 100.0%; Pred. No. 10; Matches 28; Conservative 0; Mismatches 0; Indels 0; Gaps

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ORIGIN

BASE COUNT

Search completed: September 1, 2002, 01:57:53 Job time: 10700 sec

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Result
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Maximum DB seq length: 2000300000
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Match Length
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          /cgn2_6/ptodata/1/ina/5A_CCMB.seq.*
/cgn2_6/ptodata/1/ina/5B_CCMB.seq:*
/cgn2_6/ptodata/1/ina/6A_CCMB.seq:*
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US-08-705-771 2
US-08-42-641B-1
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US-08-913-159-3
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US-08-913-159-7
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US-08-811-094-15
US-08-811-094-23
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PCT-US94-11121-23
US-08-910-3680-158
PCT-US91-03680-155
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US-09-385-982-359
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2487.850 Million cell updates/sec
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Sequence 15, Appli Sequence 1, Appli Sequence 15, Appli Sequence 17, Appli Sequence 18, Appli Sequence 185, Appli Sequence 185, Appli Sequence 185, Appli Sequence 187, Appli Sequence 188, Appli Sequence 187, Appli Sequence 187, Appli Sequence 188, Appli Sequence 188, Appli Sequence 188, Appli Sequence 188, Appli Sequence 187, Appli Sequence 188, Appli Seque
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Query Match  Best Local Similarity 100.0%; p  Matches 20, Conservative 0,  Matches 111111111111111111111111111111111111	RESULT 1  POTTOSYL USBNO-156  SEQUENCE 156, APPLICATION: APPLICANT: MATTERYK, Steven ITILE OF INVENTION: CRUSSLINKING AGE ITILE OF INVENTION: DUPLEX DNA NUMBER OF SEQUENCES: 158  COMPRESSONDENCE ADDRESS: ADDRESSEE: MOTTION & FOOTSOT STREET: 54° Middlelield Road, Suit CITY: Menlo park STATE: California COUNTRY: USA ZIP: 94025 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SUFTWAKE: PALENTIA RELEASE #1.0, V CURRENT APPLICATION DATA: APPLICATION UNMAREP PCT/USG1/03ARO FILING DATE: 19319524 CLASSIFICATION UNMAREP RECHENDE/DOCKET NUMBER: 4610-0011 TELECOMMUNICATION INFORMATION: TELEPRIONE: 415-327-250 TELEFAX: 706141 INFORMATION FOR SEQUING: SEQUENCE: CHARACIER SIGNIE STEANDEDNESS: SIGNIE TOPOLOGY: inear	2H 17 1.8 3035 2 298- 29 17 1.8 3630 3 08- 30 17 1.8 3630 3 08- 31 17 1.8 5162 2 08- 2 2 17 1.8 5162 2 08- 2 33 17 1.8 5162 2 08- 2 34 17 1.8 5162 3 08- 3 4 17 1.8 5162 3 08- 3 5 17 1.8 6769 1 08- 3 7 17 1.8 6769 1 08- 3 8 17 1.8 6769 1 08- 4 0 17 1.8 6769 1 08- 4 17 1.8 6769 1 08- 4 17 1.8 6769 5 108- 4 17 1.8 6769 5 108- 4 17 1.8 6769 5 108- 4 17 1.8 6769 5 108- 4 17 1.8 6769 5 108- 4 17 1.8 6769 5 108- 4 17 1.8 6769 5 108- 4 17 1.8 6769 5 108- 4 17 1.8 6769 5 108- 4 17 1.8 6769 5 108- 4 18 17327 1 08-
Score 20; DB 5; Length 25; Pred. No. 13; , Mismatches 0; Indels	03680  PECIFIC NONPHOTOACTIVATE NG AGENIS WHICH BIND TO  Proper 1.0, Version #1.25  /03680  0-0011.40	0S-88-883-515 1  0S-98-723-624-16 S  0S-98-74-74-000A-5  US-08-712-157-5  US-08-723-170-13 S  US-09-225-170-13 S  US-09-225-170-13 S  US-09-225-170-13 S  US-08-487-784-20 S  US-08-487-794-20 S  US-08-488-0118-20 S  US-08-488-0118-20 S  US-08-488-0118-20 S  US-08-488-0118-20 S  US-08-68-727-20 S  US-08-88-727-20 S  US-08-88-727-20 S  US-08-88-727-20 S  US-08-88-727-20 S  US-08-98-88-178-20 S  US-08-98-88-18-18-20 S
0; Caps	THE MAJOR GROOVE	GENERAL INCOMMI Sequence 18, Appli Sequence 5, Appli Sequence 13, Appli Sequence 13, Appli Sequence 13, Appli Sequence 20, Appli Sequence 21, Appli Sequence 23, Appli

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US 08 424 641B 1
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                                                                                                                                                                                                                                                     Sequence 1. Application US/884.4641B
Patent No. 5824524
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Overy Match 2.0% Score 19: D8 G Teagth 1720; Best Tacai Similarity Tooluk Pred. No. 23;
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Patent No. 6054289
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                                                                                                                                                                                                                                  GENERAL INFORMATION:
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                                                                                                                                                        APPLICANT: Sylvin Manasan, Shirley A.
APPLICANT: Walker, bessere a Valabulah ApplicaNT and sever A Valabulah di APPLICANT: and sever A Valabulah di APPLICANT is salated DNA Encoding
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 974-994-170
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                                                                                   CORRESPONDENCE ADDRESS
                                                                                                        NUMBER OF SECUENCES:
                                                                                                                        TITLE OF INVENTION:
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COUNTRY:
               STATE
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                                                                    ADDRESSEE:
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T: 2190 Commons Pareway
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; Patent No. 5925388
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Hest Local Similarity 1070.0%; Pred. No. 21;
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INFORMATION FOR SEQ ID Not 1:
                CLASSIFICATION: 435
                                                                                            CONFUTER: AFOR STEERING MS INS. STETMARE: WOLDHOUTHOUT S. ].
                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Isolated DATITLE OF INVENTION: Enzyme For TITLE OF INVENTION: Resistance
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TELEPHONE: (517) 347-4104
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NAME: Idn C. McLeod
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PHIOR APPLICATION DATA:
APPLICATION NUMBER: 08/366,480
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                                                       APPLICATION NUMBER:
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MEDIUM TYPE, storage
COMPUTER: Acer
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MEDIUM TYPE: Storage
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APPILICATION NUMBER: 08/424,641
                                                                                                                                                                                                                                                     TOUNTRY:
                                                                                                                                                                                                                                                                           STATE:
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                                                                                                                                                                                                                                                                                                        2190 Commons Parkway
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Best Local Similarity 100.0%;
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                                    TELEX: No. 5972673e
INFORMATION FOR SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb
MEDIUM TYPE: storage
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NAME: Ian C. McLeod
REGISTRATION NUMBER: 20,
SEQUENCE CHARACTERISTICS:
LENGTH: 2987 base pairs
                                                REFERENCE/DOCKET NIMFREP: QUI TELECOMMUNICATION INFORMATION: TELEPHONE: (517) 347-4100 TELEPHONE: (517) 347-4103 TELEX: No. 5972673e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sylvain Moineau, Shirley A. APPLICANT: Walker, Ebenezer R. Veddamithu, APPLICANT: and Peter A. Vandenbergh TITLE OF INVENTION: Isolated DNA Encoding
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                                                                                                                                                       ATTORNEY/AGENT INFORMATION: NAME: Ian C. McLeod
                                                                                                                                                                                        PRIOR APPLICATION DATA: 08/424, APPLICATION NUMBER: 08/424, FILING DATE: APPLI 19, 1995 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION:
TELEPHANE (517) 447-44 nin
TELEPHAX: (517) 347-44103
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                                                                                                                                       RECISTRATION NUMBER:
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                                                                                                                       Quest 4.1-155
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US-08-913-159-1
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; Patent No. 6300109
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INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER: DK 0

FILING DATE: 17-FEB-1995
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                                                                                                                                                                                                                                    FEATURE:
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                                                                                           OTHER INFORMATION:
                                                                                                                           OTHER INFORMATION.
                                                                                                                                                           IDENTIFICATION METHOD: /CO
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TYFE: nucleic acid
STRANDEDNESS: double
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STRANDEDNESS: Double
 LOCATION:
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LOCATION: 769..1620
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2412..3323
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restriction modification systems from Lactococcus lactis
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                                                                                   OD: experimental /codon_start=1613 /codon_start=1613 /product= "LIAAI -GAMC- adenine methylase H" /vevidence - EXPERIMENTAL /gene= "ODE" / Dumber - 2
                                                                                                                                                                                                                                                                                                        /product= "LlaAl -GATC- N-6-adenine methylase A" /evidence- EXPERIMENTAL / qene- "ORF"
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luû.û%; Pred. No. 21;
ive 0; Mismatches
                                                       /label- m-!laAIH
                                                                                                                                                                                                                                                     /standard_name- "Gene coding for M.I.IaAIA" /label= m-liaAIA
                                                                      /standard_name= "Gene coding for M.I.laAiB"
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                                                                Survey Match 2.5%: Severe les 98.4; Georgia 8695; Beest Local Similarity 100 9%; Pred. No. 21;
                                                     Matches
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                507 atquattqatqanagagtt 525
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STRANDEDNESS: double
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Zeodor "LlaAI GATC adenine methylase R"
Zeotok see EXPERIMENTAL
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September 7. Applib at teach 0.0/10091 0.00

Patient No. 6300109
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2 Patent No. 6400109
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HILLE OF INVENTION: restriction modification systems from factoroccus lactis
NUMBER OF SEQUENCES: 14
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AFFILING DATE: 17 FEB 1995
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CURRENT APPLICATION DATA:
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TITLE OF INVENTION: restriction modification systems from Lactococcus lactis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Lactococcus lactis subsp. cremoris
STRAIN: W9
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COMPUTER: IHM PC COMPATIBLE
OPERALING SYSTEM: PC DOSYMS DOS
                                                              SHETWARE
                                                                                                               MEDIUM TYPE: Floppy disk COMPUTER: IBM FC compatible operating SYSTEM: For the SZMS-theS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IoPoLogy: linear
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penness: double
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                                                Patentin Release #1.0, Version #1.4
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/Tabel miliaAlB
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US-(-8-378-588-15
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CEMERAL INFORMATION:
CEMERAL TOWN: John, Mallyakal E.
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Best Local S
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INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 3695 base pairs
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J.
REGISTRATION NUMBEP: 27,386
REFERENCE/DOCKET NUMBER: 111
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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APPLICATION NUMBER: DK 0179/95
FILING DATE: 17-FEB-1995
                                                                                                                                            MEDIUM TYPE: Floppy cisk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CUBRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: PRODUCING HELEKOLOGOUS PEROXIDASE
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                                                                                                         FILING DATE:
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OTHER INFORMATION:
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                                                                                            CLASSIFICATION:
                                                                                                                             APPLICATION NUMBER: US/08/378,588
                                                                                                                                                                                                                                                                         COUNTRY.
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LOCATION: 2412.
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                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Nicholas J. Seay, Quarles & Brady
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                                                                                                                                                                                                                                                        RY. USA
53701-2113
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/product = "LlaAI restriction endonuclease"
/evidence = EXPERIMENTAL
/gence = "CRF"
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; Patent No. 5869720
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Hest Local Similarity
Matches 18; Conserv
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1ELEFAX: (608) 251-9166
INFORMATION FOR SEQ ID NO: 15
                                                                                                                                                                                                  TELEFAX: (608) 251-9166
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 1064 base pairs
                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                  FILING DATE: 25-JAN-1995
ALTORNEY/AGENI INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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TITLE OF INVENTION: PRODUCING HETEROLOGOUS PEROXIDASE
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STREET:
CITY: Ma
                                                                                                                                                                                                                                                                  NAME: Seay, Nicholas J. REFERENCE/DOCKET NUMBER: 27,386
              Local Similarity
                                                                                                                   TOPOLOGY:
                                                                                                                                                    TYPE:
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
GPERALING SYSIEM: PC-DUS/MS-DOS
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STRANDEDNESS: double
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ilarity 100.0%; | Conservative 0;
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                                                                                                                   linear
                                                                                                                                                                                                                                  (608) 251-2484
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                                                                                                    CDNA
                                                                                                                                   double
              1.9%; Score 18; DB
100.0%; Pred. No. 61;
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 Mismatches
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                              DB 2;
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Sequence 15, Application 15,710; 411121
GENERAL INFORMATION:
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                                                                                                                                                                                                                                      Sequence 24. Application US/OBCASHR
Fatent No. 5608148
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                                                                                                                                                                                                                     GENERAL INFORMATION:
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TELEFAX: (608) 251 9155
INFORMATION FOR SEQ 1D NO
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NAME: SCAY, NICHOLAS
REGISTRATION NUMBER: ....
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ADDRESSEE: Nicholas' Soly, Quarles & Riady
STREET: First Wisconson Libra, one South
STREET: Finckney St.,
STREET: P.O. Box 2113
                                                                                                                        NUMBER OF SEQUENCES: ...
                                                                                                                                                              TITLE OF INVENTION: FROM DESCRIPTION PLANTS FEROXIDASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patenth Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                   APPLICANT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 1064 hase paid
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                                STREET:
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COUNTRY
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               STATE:
                                                                   STREET: First Wisconsin raid, one South STREET: Pinckney St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/LOCKET NUMBER
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Z.S.A
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INFORMATION FOR SEQ ID NO:
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MEDIUM TYPE: Floppy disk
COMPUTER: FREE SYSTEM FOR FORMS TWO
SPERATING SYSTEM FOR FORMS TWO
SPETWARE: Pated In Release #1 0, Version
FURRENI APPLICATION DATA:
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ANDRESSEE: Nicholas J. Seay, quaries & Brady
SIREEL: First witsconsit Plaza, and South
SIREEL: Picokocy St.,
SIREEL: Picokocy St.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: John, Malbyakal E.
HTHE OF INVENTION: TRANSCENIC COLLON PLANTS
THELE OF INVENTION: PRODUCTING HELEROLOGOUS PEROXIDASE
                                                                                                                    PRIOR APPLICATION DÂTA:
APPLICATION BUBBBB: 08/378/508
FILING DATE: 25 JAN 1995
ALTERNAY/AGENT INFORMATION:
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                                   REGISTRATION NUMBER: 27, 886
REFERENCEZDSCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
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CITY: Madison
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COMPUTER: THM PC COMPATIBLE
CONFECTING SYSTEM: PC DOS/MS DOS
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TYPE: nucleic acid
STRANDEENESS: double
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  TELEPHONE:
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: (608) 251 2484
(608) 251 9168
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PCT: US94-11121-23
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Search completed: September 1, 2002, 03:12:34
                                                                               υb
                                                                                                                                                                                                                                                                                                                                           NAME: Seay, Nicholas J.

REGISTRATION NUMBER: 27,386

REFERENCE/DOCKET NUMBER: 11-229-9076-8

TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 251-2484

TELEFAX: (608) 251-2166

INFORMATION FOR SEO ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 1270 base pairs

TYPE: DUCLEIC acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 23, Application PC/TUS9411121
(CEMERAL INFORMATION: John, Maliyakal E.
TITLE OF INVENTION: TRANSGENIC COTTON PLANTS
TITLE OF INVENTION: PROJECTIVE HETEROLARDIS PEROXIDASE
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                          Query Match 1.9%; Score 18; DB Best Local Similarity 100.0%; Pred. No. 60; Matches 18; Conservative 0; Mismatches
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SEQUENCE CHARACTERISTICS:
LENGTH: 1270 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 53701-2113
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPACIBLE
OPERATING SYSTEM: PC-1005/MS-1005
SOFTWARE: Patentin Release #1.0, Version #1.25
CURPENT APPLICATION NATA:
APPLICATION UNMBER: PCT/US94/11121
FILING DATE:
FILING DATE:
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                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DNA (genomic)
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CITY: Madison
STATE: WI
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TOPOLOGY: li
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Maximum DB seq length: 20000000000
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Perfect score:
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                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
  Score
Query
Match Length )B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gataacatccaatccaaccaatcacaaccaatcctgatgagataacccactttaagcccac 540
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | CCalagagtCagttaactcatttttatatttcatagatcaaataagagaaataacgqtat 420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AX105267.1 GI:13921417
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            artificial sequence.
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Best Local Similarity 100.0%; Pred. No. 1.2e-16;
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Rance, I., Theisen, M. and Gruber, V.
Patent: FR 2791358-A 2 29 SEF 2000;
MERISTEM THERAPEUTICS (FR)
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synthetic construct
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Rance, L. Theisen, M. and Gruber, V.
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MERISTEM THERAPEUTICS (FR)
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promoter MPr1096"
                                                                                                          /note "Promoter MFr1097 derived from the promoter petE by a deletion in 5' of the repeated invert sequences as well as the as-1 like box borne on fragment Sphl of 212 bp promoter MFr1097"
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1. .623
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                                                                                                                                                                                                             The pea plastocyanin promoter directs sell specific but not full light regulated expression in transferic tobacco plants plant J. 3 (3), 437-449 (1993)
                                                                                                                               AT I is a plant packed protein binding to the sequence ANTAUTITIATT in the promoters of light regulated genes (N. Datta & A.R. Cashmere 'Briding of a Bea nuclear protein to promoters of
                                                                                                                                                                                                                                                                                                               Submitted (09 SEP 199.) J. Gray, University of Cambridge, Dept. of Plant Sciences, bowning Street, Cambridge CB2 3EA, UK
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Spermatophyta: Maquoliophyta: endicutyledons: core endicots:
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/codon_start 1
/product "plastocyanin"
/protein_id "CAAAH93.1"
/db xref-"C1:437486"
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/note "importect direct repeat 1"
                                                                           /strain "Foltham tirst"
/db vrot="tavon:3888"
                            /clone lib "lambda EMBL4"
                                               /tissue_type "shoots"
                                                            /cione-"iambda GPC⊀"
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AX036607
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57; Conserv
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/product="mature plastocyanin (AA 1-99)"
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/citation=[1]
59. .88
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813. .1319
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25. .62
                                  lb-pcr"
                                                                                   /organism-"synthetic construct"
/db_xref="taxon:32630"
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ASLKDFGVALVATAASAVLASNALAVEVLLGASDGGLAFVPSSLEVSAGETIVFKNNA
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/db_xref="GI:20846"
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114. .142
                                              block oligonucleotide for the construction of promoters
                                                                 /note- Directional building block S3-Directional building
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94. .139
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100.0%; Pred. No. 9.8e-17;
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Pance, I , Theisen, M and Gruber, V.
Patent: PR 2791358-A 6 29-SEN-2000;
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liarity lim m%; Pred No. 1 5e-Conservative n; Mismatches
                                                                            reference promoter on the promoter petE. promoter Mpr1098" a 58 c 21 g 61 t
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1 67 c 31 c
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/db_xref="taxon:32630"
                                                                                                                     /hote "The promoter MP11098 of 207 bp only contains the 'TMIA' and 'CAAT' boxes and corresponds to a minimal
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100.0%;
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Logal Similarity 109.0%; Pred. No. 9.09041;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31; Conservative
                                                                                                                                                                                                                                                               Rance, I., Theisen, M. and Gruber, V. Patent: FF 2/91848 A < 29 SEP 2000;
                                                                                                                                                                                                                                                                                                                                  synthetic construct
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                                                                                                            Znote "Promoter MPrilli created by inserting at -99 bp position of MPril098, an IR by element containing a 'G' box and fusing a sequence of 58 bp (duplication of the element
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CaMV promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Znote "The promoter MPrIIIO was created by inserting at position 99 bp of MPrI098 ar element of 18 bp containing a "T" box and by fusing a sequence of 44 bp of the RNA 358
                                                                             promoter MPr}llI™
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8.7%; Score 27; DB 6; Length 7z; Best Local Similarity 100.0%; Pred. No. 0.04z; Matches 27; Conservative 0; Missurches
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Matches 41; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                 synthetic construct
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AX036609
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AX036602
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1 (bases 1 to 72)
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                                                                                                                                                           blook oligomolectide for the construction of promotors by
                                                                                                                                                                            /organism "synthetic senstruct"
/db_xref="taxon:32630"
/note="Directional building block S5 prectional building
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23: /SIDSI/gcgdata/geneseq-geneseqn-embl/NA2000.B.DAT: *
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result		Query				
70.	Score	Match	Match Length DB ID	DB	ID	Description
-	731	100.0	731	ci ci	MS02128	Promoter #3 tor re
IJ	731	100.0	971	ر <u>ا</u>	AAS02127	Promoter #2 for re
w	731	100.0	1350	د: د:	AAS02126	Promoter #1 for re
4	57	7.8	326		AAA96459	Nucleotide seguera
5	57	7.8	623	21	AAA96458	Nucleotide sequenc
6	57	7.8	834	21	AAA96457	Nucleotide sequenc
7	34	4.7	68	2	AAA96468	Nucleotide sequenc
8	34	4.7	250	1.	AAA96462	Nucleotide sequenc
9	32	4.4	207	2	AAA96460	Nucleotide seguene

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AAH70954 AAX97750	AAI 45874	AAC74357	AAA54399	AA:89182	481852	AAI89682	AAS38837	AAI83509	AAAGEERR	AAS16182	AAC 50614	AAC34680	AAH87842	M233774	AAA44596	AAA96476	ABA:8639	VISTARVED	VRF 35 300	ABL26662	ARI 12249	AAX95543	AA!59044	WIE0830	AAA96467	ABLSEETE	AAA36463	AAA96477	AAA96466	AAT77099	AAA96470	AAA96464	AAA96165	AAA96463
Human cervical can Extended human sec	rigornan	Human secreted pro	Plant defensin cod	Human polynucicoti	Human polynuolooti	Human polynucleoti	Novel human diagno	Human polynucleoti	Number tide sugardar	lumour suppressor		Arabidopsis thalia	Peppermint plant o	Tobacco plant resi	Human secreted exp	Nucleotide sequenc	Human nervous syst	Drosophila melanog	Human immune syste	Drosophila melanog	Human immune syste	wagleis acid seque	Human polynucicoti	Buman polynucleoti	Nucleotide sequenc	Human immane syste	No. location or goest	Nucleotide sequene	Karlootide sequene	Pisum sativum plas	Nucleotide sequenc	Nucleotide sequenc	Mucleotide sequenc	Nucleotide sequenc

## ALIGNMENTS

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Novel promoter for regulating expression of foreign genes in transquare organisms, more specifically in a leaf specific manner in transquare plants -	Whi; 2001-266316/27.	Vezina I., D'Aoust M;	(MEDI-) MEDICAGO INC.	04-00T-1999; 99US-0157129.	02-00T-2000; 2000WO-CA01144.	12-APR-2001.	WO200125455-A2.	Synthetic.	Promotor: 'ranegonio plant: #smoostyledon; divotyledon: tymmosperm: ds	Promoter #3 for regulating expression of foreign genes.	18-JUL-2001 (first entry)	AAS02128;	LT 1 2128 AAS02128 standard; DNA; 731 HP.

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AAS02127
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Matches 731; Conservative
AAS02127 Standard: INA; 971 (d)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organisms, particularly plants, e.g. monocotyledons, dicotyledons or
dymnosperms, by freparing a transpeal's organism using an expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The sequence represents the Solite sequence of promoter #3 for regulating expression of foreign genes of transquale organisms. The promoter is useful for regulating the expression of foreign genes in transquale
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Pred. No. 5.2e 297;
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The sequence represents the coding sequence of promoter % for regulating expression of foreign dones in transports organisms. The promoter is useful two regulating the expression to foreign decays to transport organisms, particularly pounts, e.g. monocodyled as, decayledges on approximation of transported organisms, by preparing a transporter organism spin at expression construct comprising the promoter and an open reading frame (CFF) of a
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481 cutttqattadacatqtqattatttaafqaattqatqaaqqaqtttqqattaaaqffqtat 94:
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                                        useful for regulating the expression of foreign genes in transgenic organisms, particularly plants, e.g. monocuty indons, dicutyledons of gymnosperms, by preparity a transgenic organism using an expression construct comprising the promoter and an open reading frame (OPE) of
                                                                                               The sequence represents the coding sequence of promoter #1 for regulating expression of foreign genes in transgenic organisms. The promoter is
                                                                                                                                         Claim 1; Page 8; 9pp; Erglish
                                                                                                                                                                                   organisms, more specifically in a leaf-specific manner in transgenic
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BP; 482
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Α,
236 C; 230
G; 402 %; 0 other;
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Query Match Best Local Similarity

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Score 731 Pred. No.

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DB 22; Length 1350; .9e-297;

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              WC200056906-A1
                                           Pisum sativum
                                                                                                                                Nucleotide sequence of the promoter MPr1096
                                                                                                                                                                                                                           AAA96459 standard: DNA; 326
                                                                                      plastocyanin dene promoter;
                                                                                                    petE promoter; chimeric promoter: transgenic plant, MPr1096
                                                                                                                                                                 08-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                           - CaddadCCddtnnanatnfffafnannnattotataaaaatoacantttgtgagtintac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   geactitytyyeaeatetaeattatetaaateacaeattetti,asacatetyayeeaca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gcactotgtggcacatotacattatotaaatoacacattetteracacatotgageeaca
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         attaatccctccaaaaaaaaaaacggtatatttactaaaaaaatrtaagccacgtaggag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Matches 57; Conservative
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Rance 1, Gruber V,
                                    (MERI ) MERISTEM THERAPEUTIC
                                                                               22 MAR 1999;
                                                                                                                     20 MAR 2000; 2000 WO (HUSS)?
                                                                                                                                                              28 SEP 2000
                                                                                                                                                                                                     Mo500028408 VI
                                                                                                                                                                                                                                             Pisum sativum
                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                     plastocyanin gene promoter
                                                                                                                                                                                                                                                                                                                           petE promoter; chimeric prometer; transgenic plant: MPr1097;
                                                                                                                                                                                                                                                                                                                                                                 Nucleotide sequence of the promoter MP:1097
                                                                                                                                                                                                                                                                                                                                                                                                        OB FEB 2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA96.458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA96458 standard; DNA; 523 of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 326 BP, 111 A; 91 C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 site. The chimeric promoters are used in expression electors for producing transqueric plants, such as drouty-lodomous species, e.g. potato, tokero, control, lottings, tomato, motors operator, pea, rape, bectroot, or sunflower, and monocotylescens; species, e.g. wheat,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2: Page 67, 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chimetre expression promoter for producing diestyledonous and monocotyledonous transquire plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             519 agataaceeaetttaagee9aeg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    barley, oat, rice, or com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WP1: 2000-587667/55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MERI ) MERISTEM THERAPLULIC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          110 agatadoreaetttaagoreaeg setetgtggoveatetaeattatetaadtegea 166
                                                                               55 35 0000 3L466
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The Useri M
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Pred. No. 1.7e 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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Best Local Similarity los.
Techos 57; Conservative
                                                                                   derived from a promoter of the pea plastocyanic tene
                                                                                                  Chimeric expression promoter for producing dicotytedates and monocofyledonous transfering plants comprises a nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                                            ₩0200056906·A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric expression promoter comprision a promoter of the pad plastromanic area, or comprision a 5 kmx operably of functionally linked upstrom of a 7AX box. That box and transcription initiation site, the chimeric promoters are used in expression vectors to product transcription in species, e.g. polaro, tobacco, cotton between tomates dicotyledomous species, e.g. polaro, tobacco, cotton between tomates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents the chimeric promoter Mp.1097. The promoter is derived from the petE promoter from post-bastocyanin dense, by a deletion in 5% of the repeated invert sequences as well as the as I like box borne on a 5th framment of 212 bp. The petE promoter directs cell specific but not tall light required expression in
                                               Claim 1; Page 65; 83pp; English
                                                                                                                                                                WPI: 2000 587667/55
                                                                                                                                                                                                                                         (MERL ) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                  20-MAR 2000; 2000WO IB00317
                                                                                                                                                                                                                                                                                                                                                                                                                                 Pisum sativum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plastocyanin dene promoter; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     petE promoter: chimeric promoter: 'ransqenic plan'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleatide sequence of the petE promoter from plast ejanin gen-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08 FEB 2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA96457 standard; DNA; 834 RP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 624 BP; 230 A; 128 P; 42 B; 183 1; 5 sather
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric expression promoter to producing dicotyledopous and monocotyledopous transquire plants comprises a purior reid sequence derived from a promoter of the pea plastocyanin scene
                                                                                                                                                                                                                                                                             22-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                        28-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        species, e.g. wheat, batiey, pat, rice, or corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPT: 2000 587667/95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  407 agataacceaetttaageecaegeaetetgtggeacatetaeattatetaaateaea 463
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                                                                                                                                                                                                     Gruber V.
                                                                                                                                                                                                                                                                               99FR 0003635
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c Pred. No. 1.6e 14.
u; Mismatches ()
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The present sequence represents the petE promoter from the pear plastoryanin gene. The promoter directs reliespecific but not

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transcription initiation site. The chimeric promoters are used in expression vectors for froducing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, contron, lettice, tomato, melon, cucumber, pea, rape, bectiout, or sunflower, and monocotyledonous species, e.g. wheat, barley, out, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            promoter of the pea plastocyanin gene, or comprising a G bex operably or functionally linked upstream of a CAAN box, TATA box and
                                                                                                                                    which is used to construct chimeric promoters of the invention in PCR reactions. The specification describes a chimeric expression promoter comprising a petE promoter of the pea plastocyanin gene, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 light-regulated expression in transpendy tobacco plants. The promoter is used to construct chimeric promoters of the invention. The
                                  and monocotyledonous species, e.g. wheat, harley, cat.
                                               plants, such as dicetyledonous species, e.g. polato, tobacco, control
lettuce, tomato, melon, cucumber, pea, rape, beetroot, or sunflower,
                                                                                   promoters are used in expression vectors for producing transquaic
                                                                                                    comprising a G box operably or functionally linked upstream of CAAT box, TATA box and transcription initiation site. The chim
                                                                                                                                                                                                 The
                                                                                                                                                                                                                               Claim 24; Page 73; 83pp; English.
                                                                                                                                                                                                                                                                   derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                  Chimeric expression promoter for producing dicotyledomous and monocotyledomous transgenic plants comprises a nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                             Rance I,
                                                                                                                                                                                                                                                                                                                                                                                                                  (MERI-) MERISTEM THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-MAR-2000; 2000WO-IB08317
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              petE promoter; chimeric promoter; transgenie plant; MPr1108;
plastocyanin gene promoter; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleotide sequence of the directional building block $3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 834 BP; 289 A; 175 C; 143 G; 237 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            specification describes a chimeric expression promoter comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        519 agalaacccactttaagcccacgcactclglggcacatctacattatctaaatcaca 575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7
                                                                                                                                                                                             present sequence represents the directional building block
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       agataacccactttaageccacgcactctgtggcacatctacattatctaaatcaca
                                                                                                                                                                                                                                                                                                                                           2000-587667/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                             Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                        The chimeric
                                    rice,
                                                                and ton,
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Query Match

Score

34;

DB

21;

Length

Sequence 68 BP; 22 A; 19 C; 5 C;

22 T; 0 other,

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AAA96462
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                                                                                                               Matches
                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 100 Matches 34: Conservative
                                                                                                                                                                                                                                                                                                                                                                                                             but not full light-regulated expression in transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric expression promoter comprising a promoter of the pea plast-cyanin gene, or comprising a 6-bex operably or functionally linked upstream of a CAAT box, TATA box and transcription initiation site. The chimeric promoters are used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    promoter is derived from the petE promoter from pea plastocyanin gene, by fusing the petE as: 1 like and nos enhancer like elements to the promoter Mpr1098 (comprising TATA and CAAT boxes of petE), and then deleting 33 bp from the 5' HTP. The petE promoter directs cell-specific
                                                                                                                                                                                                                                                                                                                              dicotyledonous species, e.g. potato, tobacco, cotton, melon, cucumber, pea, rape, beetroot, or sunflower, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chimedic expression promoter for producing dicotyledonous armonocotyledonous transgenic plants comprises a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA96462 standard; DNA; 250
                                                                                                                                                                                                                                                  audentes
                                                                                                                                                                                                                                                                                                    species, e.g. wheat, barley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                      expression vectors for producing transgenic plants, such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents the chimeric promoter Mpr1109.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2, Page 69, 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      derived from a promoter of the pea plastocyanis gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rance I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-MAP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-MAR-2000; 2000WQ-1H00317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₩02/nijinskatib -A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pisum sativum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    plastocyanın dene promoter; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 petE promoter; chimeric promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleutide sequence of the promoter MPr1109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-164-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA96462;
                          542 cactctgtggcacatctacattatctaaaatcaca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 542 cactetytyycacatetacattatetaaateaca 575
                                                                                                                                         Local Similarity
91 cactetytyycacatotacattatotaaatgaca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12 cactetytygcacatetacattatetaaateara 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8
                                                                                                                                                                                                                                                     250 BP; 81 A; 67 C; 31 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gruber V,
                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99EB-0003635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; F
                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Theisen M
                                                                                                                                                                4.78;
                                                                                                               0.
                                                                                                                                         Pred No 7 60-0
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Mismatches 0;
                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transgenic plant; MPr1109
                                                                                                                                                                                                                                                     /1 T; 0 other;
                                                                                                                                            7 60-05
                                                                                                                                                                   Length 250;
                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Slebui
                                                                                                                                                                                                                                                                                                                                 and monocutyledonous
                                                                                                                                                                                                                                                                                                                                                              lettuce, tomato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seduence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ..
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AAA96460
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                                                                                                                                                                                    KESSELF 10
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Best Local Similarity 100.0%;
Matches 42; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transgenic tobacco plants: The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operatory or functionally linked upstream of a CAAF box. TAFA box and transcription initiation sets. The chimeric promoters are used in expression well as to producing transcence plants, such as dicetyledamons species, e.g. potato, tobacco, cotton, lettuce, tomato.
petE promoter; chimeric promoter; transgenic plant; MPrIII0
                                Nucleotide sequence of the promoter MPT1110
                                                                        08 FEB 2001
                                                                                                                                               AAA96463 Standard; DNA; 280 op
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 207 BP; 67 A; '88 C; 21 G; 61 I; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                       meton, cucumber, pea, tape, bectroot, or suntlower, and monocotyledonous species, e.g. wheat, barley, ed., tice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      minimal reference promoter on the petE promoter. The petE promoter directs cell specific but not full light regulated expression in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents the chimeric promoter Mpr1098. The promoter is derived from the petE promoter from pea plastocyanin gene, and contains only the TATA and CAAT boxes, and corresponds to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; Page 67, 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chimeric expression promoter for producing d cytylodogogs and monocotyledonous transaccic plants comprises a nocleic acid sequence derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPT: 2000 587667755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rance I, Gruber V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
Pisum sativum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       plastocyania deno promotor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MERI ) MERISTEM THERAFERITOS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0007 dBS 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W0200056906 AL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleotide sequence of the promoter MPr1098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA96460;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA96460 standard; DNA; 207 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22 MAR 1999; 99FR 000 6.35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       petE promoter; chimeric promoter; transgenic plant. MPr1097;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08 FEB 2001 (lirst entry)
                                                                                                                                                                                                                                                         544 etetqtqqeaeatetaeatrateraaateaea 875
                                                                                                                                                                                                                                           16 etertytygeaeatetaeattateraaateaea 47
                                                                        (tirst entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The issent Ma
                                                                                                                                                                                                                                                                                                                             Score 32: DB 21; Leigth 207;
Pred. No. 0.00053;
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Best Logal Similarity 100.0%:
Watches 41: Conservative 0
                                                                                                                                                                                                                                                                                                                           AAA96465 standard; DNA; 296 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 280 BP; 88 A; 73 C; 42 G; 77 T; 0 other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            box, TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transquaic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, bectroot, or sunflower, and monocotyledonous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
                  28-SEP 2000
                                                        WO200056906-A1
                                                                                              Pisum sativum
                                                                                                                                                  plastocyanin gene promoter; ss
                                                                                                                                                                                                          Nucleotide sequence of the promoter MPrilli.
                                                                                                                                                                                                                                              08-FEB-2001 (lirst entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inserting a 6 bex and lucing a tragment of 350 kMA of CMV. The potE promoter directs cell-specific but not tall light regulated expression in transagente tobacco plants. The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents the chimeric promoter Mprillo. The promoter is derived from the petE promoter from the petE promoter from the petE promoter from the petE promoter that clements to the by tasing the petE as I like and now cohancer like clements to the promoter MPrilo88 (comprising TATA and CAAT boxes of petE), and then
                                                                                                                                                                    petE promoter: chimerie promoter: transdenie plant; MPrill!;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                species, e.g. wheat, barley, oat, rice, or corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operably or functionally linked distream of a CAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2; Page 70; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22 MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plastocyanin gene promoter; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                           542 caetetgtggeaeatetaeättatetaaate 572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MERI ) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20:MAR-2000; 2000WO-TB00317
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                                                                                                                                                                                                                                                                                                                                                                                                                       68 cactetqtqqcacatetacartatetaaate 98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.2%; Score 41; DB 21; Length 280;
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O; Mismätches J
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В
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Best Local .
                                                                                                                                                                                               Pisum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inserting a G box and fusing a fragment comprising a duplication of the element as2 and as1. The petE promoter directs cell-specific but not full light-regulated expression in transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operably or functionally linked ubstream of a CAAT box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              promoter is derived from the petE promoter from pea plastocyanin gene, by fusing the petE as-1 like and nos enhancer like elements to the promoter MPr1098 (comprising TATA and CAAT baxes of petE), and then
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        monocotyledonous transgenic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene \,
                                                                                                      20-MAR-2000; 2000WO-IB00317
                                                                                                                                      28-SEP-2000
                                                                                                                                                                   W0200056906-A1
                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                             plastocyanin
                                                                                                                                                                                                                                                                                         Nucleotide sequence of the promoter MPr1153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 296 BP; 94 A; 74 C; 45 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents the chimeric promoter Mprllll. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Fage 71; 83pp, English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chimeric expression promoter for producing dicotyledonous and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rance I, Gruber V,
                Rance I, Gruber V,
                                            (MERI-) MERISTEM THERAPEUTICS
                                                                           22-MAR-1999;
                                                                                                                                                                                                                                                            petE promoter;
                                                                                                                                                                                                                                                                                                                       08-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                    AAA96464 standard; DNA; 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                barley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-MAR 2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MERI-) MERISTEM THERAPETITICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             542 cactotigtigcacatotacattatotaaato 572
                                                                                                                                                                                                                                                                                                                                                                                                                                                              84 cactctgtggcacatctacattatctaaatc 1]4
                                                                                                                                                                                                                                                                                                                                                                                                                   12
                                                                                                                                                                                                  sativum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               beetroot, or sunflower, and monocotyledonous species, e.g. wheat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                             gene
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                                                                           99FR-0003535
                                                                                                                                                                                                                                                          chimeric promoter; transgenic plant; Mar1153;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99FR-0003535
                                                                                                                                                                                                                                             promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                Theisen M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    83 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.0014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 296
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AAA96470
ID AAA9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
          The present sequence represents the directional building block swhich is used to construct chimeric promoters of the invention in PCR reactions. The specification describes a chimeric expression
                                                                                                                         Chimeric expression promoter for producing directyledonous and monocotyledonous transgenic plants comprises a nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          construct chimeric promoters of the invention. The specification describes a chimeric expression promoter domprising a promoter of the pea plastocyanin gene, or comprising a 6 box operably or functionally linked upstream of a CANT box, TATA box and transcription initiation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Page 70: 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nucleic acid sequence derived from a promoter of the pea plastocyanin gene
                                                                             Claim 24,
                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sunflower, and monocotyledonous species, e.g. wheat, bariey, oat, rice,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cutton,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents the chineric promoter Mpril53. The promoter is derived from the petE promoter from pea plastocyanin gene. The petE promoter directs cell-specific but not full light regulated expression in transferic tobacco plants. The promoter is used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI, 2000 587667/55
                                                                                                           derived from a promoter of the pea plastocyanin gene
                                                                                                                                                                                                            Rance 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                             plastocyanin gene promoter; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleotide sequence of the directional building block S5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA96470;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA96470 standard, DNA;
                                                                                                                                                                                                                                             (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                              22-MAR-1999;
                                                                                                                                                                                                                                                                                                             20-MAR-2000;
                                                                                                                                                                                                                                                                                                                                              28-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                           petE promoter; chimeric promoter; transgenic plant; MPrll08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            542 cactctgtggcacatctacattatctaaatc 572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91 cactetgtggcacatetacattatetaaate 121
                                                                                                                                                                              2000-587667/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              lettude, tomato, meion,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              सार RP; lul A; 7x ते; 41 ते; सर ग्रं u other;
                                                                            Page 75; 83pp,
                                                                                                                                                                                                            Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                             2000WO-IB00317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                            English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 31;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              eucumber, pea, rape, beetroot,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.0014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 303;
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C.

promoter comprising a petE promoter of the pea plastocyanin gene, or

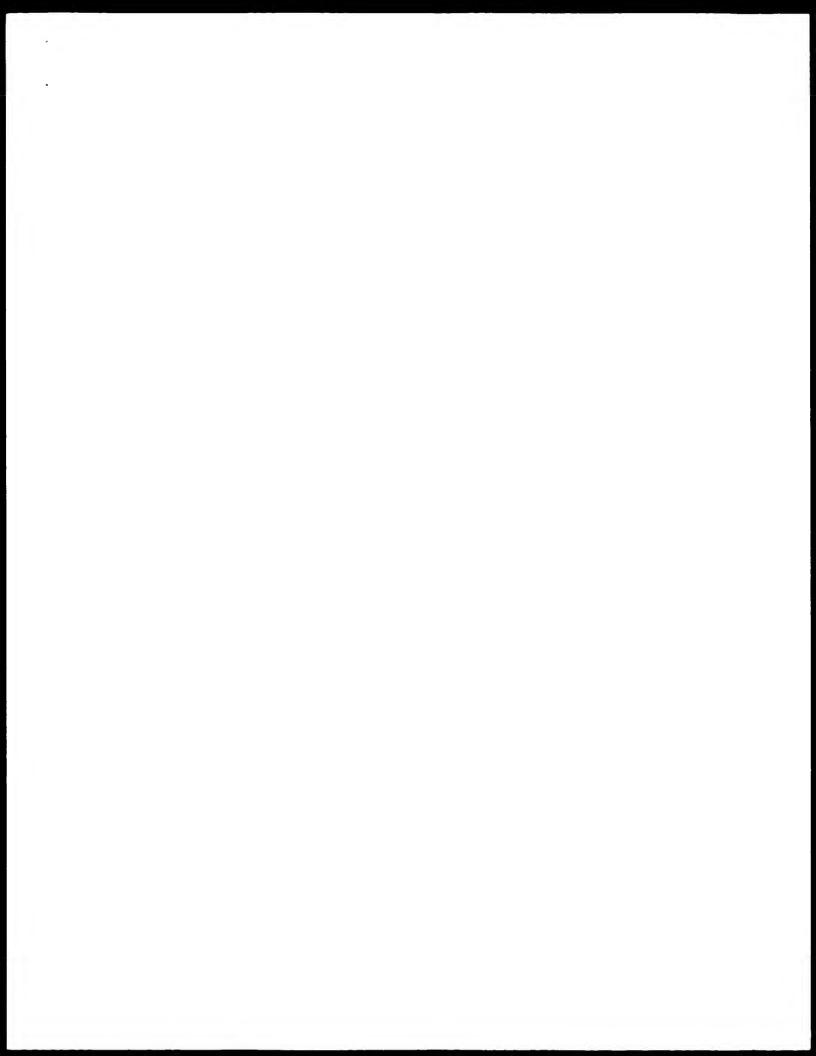
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                         the GUS reporter dens in tobacks where the GUS dens was under the control of the PetE promoter and 458 CaMV promoter. The enhancer was also used joined to the (48) to all region of the patalin promoter PS20
                                                                                                                                                                                               This sequence represents the pea plastocyanin promoter enhancer corresponding to nucleotides 444 to 179 of the full length sequence. This enhancer has been used to increase the expression of
                                                                    which was joined to the 70% reporter and transformed into potato using Altimetacieus. Expression of the december of one of more genes of a plant increases in green or non-green tissues, especially roots, tubers, seeds, stems, flowers of leaves. This enhancer may operate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comprising a G box operably of lunctionally linked upstream of a CAAT box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic
Sequence 266 BP; 102 A; 42 C
                                    both in normal and reverse directions and can be attached either to
the promoter of terminator of the gene to be expressed.
                                                                                                                                                                                                                                                                           Claim 9; Figure 6; 46pp: English
                                                                                                                                                                                                                                                                                                            Enhancer from pea plastoryan's promoter region—used to produc
transagenic plants having increased expression of a desired gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         promoter
                                                                                                                                                                                                                                                                                                                                                                   Mb1: 1887 310010728
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              × 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pison sativon plastocyano permoter region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAI77099 standard: DNA; 255 m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, meton, cucumber, pea, rape, bectroot, or sunflower, and monocotyledonous species—e.g. wheat, barley, oil, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29 NOV 1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26 NOV 1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       W09720056 A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  plastocyanin; promoter; enhancer; reporter gene; per;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 JAN 1998 (trist early)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 72 BP: 22 A; 20 C; (4 G; 20 I; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        546 etgtggeaeatetaeattatetaaate 5/2
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 34 G; 83 L; 0 other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                             used to produce
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AAA96466
                                                                      Best Local Similarity
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Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                            by fusing the petE as:1 like and nos enhancer like elements to the promoter MFr1098 (comprising TATA and CAAT boxes of petE), and then inserting a 6 box and fusing a fragment comprising a duptication of the element as2 and as1. A 72 by fragment comprising the classics at 2, as and as 1 was then deleted. The petF promoter directs cell specific but not full light-regulated expression in transquare totacee plants, the promoter is used to construct clauser promoters of the invention. The specification describes a chimeric promoters of the invention.
                                                                                                                                                                                                                                                               e.q. potato,
                                                                                                                                                                                                                                                                                                          promoter of the pea plastocyanin wene, or comprising a 6 hox operably or functionally linked upstream of a CAAL box, IALA box and transcription initiation site. The chimeric promoters are used in expression vectors
                                                                                                                                                                       Sequence 220 BP: 73 A; 60 C; 26 G; 6] I; 0.3(E)-r
                                                                                                                                                                                                                  barrey, out, rice, or corn
                                                                                                                                                                                                                                           rape, beetroot, or suntlower, and monocotylesbosous species, e.g. wheat
                                                                                                                                                                                                                                                             tot production transactic plants, such as dicorpletionous species, e.g. putato, tobacco, collon, lettuce, tomato, melon, enemiher,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents the chimeric promoter Mpril143. The promoter is derived from the petE promoter from psa plastocyanin deno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2; Page 72; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    derived from a promoter of the pea plustocyanin dene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric expression promoter for producing dicotyledomous and monocotyledomous transgenic plants comprises a nucleur acid sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MERI ) MERISIEM THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plastoryanin dene promoter; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               petE promoter; chimeric promoter: transgenic plant: MFril4 o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleatide sequence of the promoter MPr1143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (Attualist) 1007 Had 80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22 MAR 1999;
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                                                                      3.4%: Score 25: [08:21]; Length 21::
190.0%: Fred. No. 0.45:
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Search completed: September 1, 2002, 03:16:38 Job time: 8287 sec



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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution

#### SUMMARIES

Result No.	Score	Query Match	Query Match Length	DB	ID	Description
c 1	39	5.3	376	1 :	HE635678	Geografia 8292898
N	ر د	œ	6:1	<u>_</u>	B1271179	BI271179 NF050C10F
ω	29	4.0	17.70	٥	AW692408	AWF92408 NPOSOHIOS
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7	29	4.0	483	10	BE249400	BE249400 NF014F06L
œ	29	4.0	603	٥	AW692904	AW692904 NF0606128
9	29	4.0	611	10	BE249469	BE249469 NF020B11L
10	47	4.∪	637	=	BG4575n3	BG457563 NF105G07P
11	29	4.0	645	10	BE249343	BE249343 NF014CC9L
12	29	4.0	683	10	BF637150	
13	28	3.8	223	10	BF518736	HF513736 EST456190
14	28	ω π	27.5	c.	AWNK3525	AWSBAS26 NEO16G101
c 15	28	3.8	468	12	AQ841643	AQ841643 T137356b
16	27	3.7	340	10	BE316138	PE316138 NF032G07L
17	27	3.7	514	10	HCC5x54x3	FG585489 EST488057

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## ALIGNMENTS

	source	FEATURES						COMMENT	JOURNAL		TITLE		AUTHORS				ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESS ON	DEFINITION	LOCUS	R⊮SULT 1 н⊬635678/с
/oranism-Medicago truncatula" /db_aret="taxon.3880" /clone_"NF109F09DI" /clone_lib="brought" /closuc_tipe="Plantlets" /fissuc_tipe="Plantlets" /dev_stage="Pooled timepoints" /hote="Vector, Lambda Zap; Contains a mixture of entire-plantlets harvested in a series of days-post watering		Seq printer: ICACACAGGAAACAGGTATGAC. iJocation/Qualifiers	Flate: 199 row: Fleedumo: 09	Email: pdmay/noble.org	F3x: 580 221 7391	2510 eam Noble Darkway, Ardmere, OK 73402, USA	le Foundat	Plant Biology Division	Unpublished (2000)	Medicago truncatula drought library	Expressed Sequence Tags from the Samuel Roberts Noble Foundation	Flores.H.B. Inman J.T. Weller J.W. and May G.D.	Torrer Jerez I. Scott A.D. Harris A.B. Gonzales R.A. Reli C.J.	Medicago.	Rosidae; eurosids I; Fabales; Fabaceāe; Papilionoideae; Tritolieae;	Eukaryota, Viridiplasiae, Sireptophyta, Embryophyta, Trachosphyta,	Medicago truncatula	barrel medic.	EST:	BF635678.1 GI:11899836	5', mRNA sequence.	DT1F1077 Drought Medicago truncatul	BF635678 376 bp mRNA linear EST 19-DEC-2000	

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Expressed Sepence lags from the Samuel Kaberts Noble Foundation
Medicago transatula flower library
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Adev stake "Seveloping tally pooled, Contains a mixture of very years, developing, fully spened flowers and flowers in early transition into pods."
Anotee "Vector: Lambda Zap: chNA was prepared from polyArenriched, pooled samples of equivalent amounts of total RNA from very years, developing, fully-opened flowers and flowers transitioning into pods. The cDNA was directionally ideated into the Unitary RNA vector.
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                              Email: gdmay@noble.org
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Expressed Sequence Tags from the Samuel Roberts Noble Foundation Medicago truncatula flower library
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                                                                                                                                                                                                      2510 Sam Noble Parkway, Ardmore, OK 73402, USA
Tel: 580-221-7391
Fax: 580-221-7380
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Insert Length: 378 Std Error: 0.0
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The Samuel Roberts Noble Foundation
2510 Sam Noble Parkway, Ardmore. OK 71402, USA
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XII-Blue MRF* (Stratagene). Excised plasmids were plated using SOLR cells."
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         29; Conservative
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2510 Sam Noble Farkway, Ardmore, GK 78402, USA
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,C.J., Flores,H.R., Imman, L.L., Weller,J.W., May,G.D., and Dixon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NF105G07PL1F1054 Phosphate starved leaf Medicago truncatula cDNA clone NF105G07PL 5', mRNA sequence.
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The Samuel Roberts Noble Foundation
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On Jul 13, 2000 th
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                    Unpublished (2000)
                                                                                                                                                                                                                                                                                  Liu,J., Scott,A.D. Harris,A.R. Genzales,F.A., Rell,C.L., Fictos, H.R., Inman,T.T., Weller,J.W., May,G.D. and Harrison,M.J. Expressed Sequence Tags from the Samuel Roberts Noble Foundation
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae, eurosids I; Fahales, Fabaceae; Fapilionoideae; Tritolieae;
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Fax: 580 221 7380
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: gdmay@noble.org
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                                                                                                                                Samuel Roberts Noble Foundation
Sam Noble Parkway, Ardmore, OK 73407, HSA
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/dev_stage "Pooled developmental"
/dev_stage "Pooled developmental"
/notea="Vector. Lambda Zap, Contains a mixture of very
young, developing, mature and senescing leaves "
138 c 132 q 177 t
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/tissue_twos="loaf"
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Insert Length: 645 Std Error: 0.00
Plate: 014 row: C column: 09
Seg primer: TCACACAGGAAACAGCTATGAC.
                                                                                                                                                                                                                                                                                                                                                                                                 2510 Sam Noble Parkway, Ardmore, GK 73492, USA
Tel: 580 221 7391
Fax: 580 221 7380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (2000)
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The Samuel Roberts Noble Foundation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: May GD
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                                                                                                                                                                                                                                                                                                                                                                           Email: gdmay@noble.orq
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                      175
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               /dev_stage="Pooled developmental"
/goote="Vector: Lambda Zap, Soutains a mixture of very
young, developing, mature and senescing leaves."
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                                                                                                              /clone_lib="Developing leaf"
/tissue_type="leaf"
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Query Match

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Enkaryota: Viridiplum no Streptophytu: Embryophytu: Tracheophytu:
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The Samme, Boberts Noble Foundation
2510 Sam Noble Firkwit. Archnore, OK 7(402, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Flores, H.R., Inman, J.E., Weller, J.W. and May, G.D. Expressed Sequence Fais from the Sammel Roberts Noble Foundation Medicase transatsia jeaf library
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Spermatophyta: Monoclophyta: endicotyledons: core endicots:
Rosidae: eurosids l: cabales: Fabaceae: Papillonoideae; Tritolicae:
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                                                                                                BES18746_1 GI:1140/474
                                                                                                                                  Sedimentes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Torres Jerez.I.,
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a. 152 c. 188 q. 204 t.
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/clone lib "leveloping leaf"
/tissue type "leaf"
/dev_stage "pooled developmentat"
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                                      AUTHORS
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679 cacatacaaaqaqaaqaqactaattaat 706
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Torres Jerez, L., Séott, A.D., Harris, A.E., Donzales, R.A., Belly'. Plores, H.E., Imman, J.T., Weller, J.W. and May, 5.D. Expressed Sequence Tags from the Samuel Robert's Noble Poundation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NEULEGIOLFIFIGES Locar-Espaine fost Medicades francestála ciema croso-
NEOLEGIOLF 57, mENA sequence
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Tel: 612-625-1243
Fax: 651-649-5058
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                                                                                                                                                                                                                          Bukuryota: Viridip antae: Streptophyta: Embryophyta: Tracheophyta
Spermatophyta: Mannoliophyta: eadicotyledoms: pose endicots;
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                                                                                                                                                                                       Rosidae: eurosids I: Fabales: Pabaceae: Papitibanolideae: Initolieae
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University of Mitthesofa Jame: M276[596
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University of Minnesota
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rosidae: eurosids I: Fabales: Pabaceae: Papillone-deae: Fritelieue
Medicago:
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                                                                                                                     (bases 1 to 359)
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56 c 44 q 57 t
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/db xref "taxon: 8880"
/clone="ptoSIL 17H17"
/clone=lib="DSI"
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/lab_host "E. coli strain XLoLP"
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                                                                          Gonzales, R.A., Bell, C.S.
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AQ841643/c
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28; Conservative
                                                                                                                                                                                                                                                         The Crop Biotechnology Center
Texas A&M University
Department of Plant Fathology and Microbiology, km 120 L.F.
Peterson Bidg, Cullede Station, TX 77843-2132, USA
Tel: 409 845 8743
Fax: 409 862 4790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A0841643 468 bp DNA librar GSS 27-SEP-1999
1937856 shotgun sub-library of BAC clone 72F02 Medicago truncatu.a
genomic clone 72FF02 084, DNA sequence.
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Insert Length: 718 Std Error: 0.0
Flate: 016 row 3 column: 10
Seq primer: TCACACACAGAAACAGCTATGAC
                                                                                                                                                                Email: dcook@ppserver.tamu.edu Other name: BSC-25-84, date. 7/2/99, Submitted to the Database of Genome Survey Sequences (GSS) on 09/27/99; More information is available at 'http://chrysie.tamu.edu/medicage'.
                                                                                                                                                                                                                                                                                                                                                                                                                       l (bases 1 to 468)
Kim,D., Choi,H., Peng,H., Ellis,L. and Cook,DR.
HAC survey sequencing of Medicago truncatula (1999c)
Unpublished (1999)
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2510 Sam Noble Parkway, Ardmore, OK 74402, USA
Tel: 580 221 7401
Fax: 580 221 7380
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On Apr 14, 2000 th
                                                                                                                                 Seq primer: pUC-C
Class: BAC subclone.
                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Cook DR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bukaryota: Viridiplantae; Streptophyta; Embryophyta; Trachcophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae;
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/clone_lib="Developing leaf"
/tissue_type="leaf"
/dev_stage="Pooled developmental"
/note="Vector: Lambda Zap: Contains a mixture of very
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  young, developing, mature and senescing leaves."
. 85 c 75 g 101 t
                                        /organism-"Medicago truncatula"
/cultivar-"genotype Al7"
/db_xre1-"taxon:3880"
/clone-"72-F-02-084"
                                                                                                       Location/Qualifiers
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    Mismatches

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Curry Match

3 8%, Source 28; PB (2) Length 468;

Best Local Similarity 100.0%; Pred. No. 8.8,

Matches 28; Conservative 0; Mismatches 0; Indels 0;

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BASE COUNT

Search completed: September 1, 2002, 01:57:56 Job time: 10703 see

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Title:
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Post-processing: Listing first 45 summaries

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Fred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result	Score	Query Match	Length	DB	ID	Description
1	19	ر دا	1720	. ب	ms-08-705-771-2	Sequence 2, Appli
۲3	19	2.6	2987	_	·08·424·6418	<del>,</del>
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	18		1884	_	US-07-704-288C·1	Sequence 1, Appli
c 17	18	-	1884	٦	пѕ-08-379-259-1	1,
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                                                                                           GENERAL INFORMATIONS
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INFORMATION FOR SECULD NO. 1
SECULENCE CHARACTERISTICS
LENGTH: 2987 base pairs
            APPELICANT: Sylvain Moisean, shirley A. APPELICANT: Walker, Element E. Vedamoths, APPELICANT: and February E. Vanfemborgh (TILLE OF INVENTION: 18 Jarot DNA Elementary)
                                                                                                                                                                                                                                     1446 ADGAALDGATGAAAGAGIT 1463
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SOFTWARE: WORTHOUT OF I
CHREENT ADDLICATION DATA:
ADDLICATION NUMBER: SS/OB
FILLING DATE: ADDLI TO, 119
TITLE OF INVENTION: Engyme out Phage
                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Lan C. McLood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FIFLE OF INVENTION: Enzyme (i) Phage FIFLE OF INVENTION: Resistance NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Sylvain Moineau, Shirley A. APPLICANT: Walker, Ebenezer R. Vedamuthu, APPLICANT: and Peter A. Van Jembergh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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CLASSIFICATION: 4(5)
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                                                                                                                                                                                                                                                                                                                                  / Match 2 + % State (9) (68 f) Langth 2987; Local Studianty 10 ok; Pred. No. 16;
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MEDIUM TYPE: Diskette S., S. modb, 080 Kb
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(517) 347 4103
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                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                Sequence 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Outry Match 2.6%: Sected by 18-2: Lebeth 2002; Best Local Similarity 100.0%: Prod. No. 16: Matches 19: Conservative 0, Mismatches of Indees
                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLEMANT: Sylvato Mothero, Shirley A.
APPLICANT: Walker, Element R. Vedamutho,
APPLICANT: and Poter A. Vandenheidh
TITLE of INVENTION: Isolated DNA Encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                COMPUTER READABLE PERMI
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ALLOWNEY/AGINE INFERMATION:
NAME: Tab C. McLood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMEDIUM TYPE: Diskette, 5.25 inch, 460 Kb
MEDIUM TYPE: Storage
                                                                                                                                                                      CORRESPONDENCE ADDRESS
                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                         TIBLE OF INVENTION: Kesistance
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      267 athdatthathdadadadtt 285
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            MEDIUM TYPE: Diskotto, 5,25 inch, 60 Kb
MEDIUM TYPE: Storado
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FILING DATE: April 19, 1995
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SOFTWARF: Wordportect 5.1
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                                                                                                                                      STREET:
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OPERATING SYSTEM: MS-DOS

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TELEPHONE: (517) 347-4100
TELEFAX: (517) 347-4103
TELEX: No. 5972673e
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                           FILING DATE: 17-FEB-19
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                   MOLECULE TYPE: DN
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                          FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 0179/95
                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.30 (EPD) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1446 ATGAATTGATGAAAGAGTT 1464
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MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES.
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||||||||||||||||||||||||
                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                         STRANDEDNESS: double
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FILING DATE: April 19, 1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE:
                                                                 STRAIN:
                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER .
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              NAME/KEY: CDS
LOCATION: 769.
                                                                                     ORGANISM
                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                            LENGTH:
IDENTIFICATION METHOD: experimental
                                                                                                                                                                                                         nucleic acid
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                                                                                                                                                                                                                          3695 base pairs
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100 0%; Pred. No 16;
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                                                                                  lattis subsp. cremoris
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Palentin Pelease #1 0, Version #1
                               ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                          MOLEGUE LATE ON CONTROL OF THE CONTR
                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 3695 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
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      ORGANISM:
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Lactoroccus lactis subsp. cremoris
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/product= "ITANI restriction endonuclease"
/evidence= EXPERIMENTAL
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/product= "LlaAI -GAIC- N-6-adenine methylase
/evidence= EXPERIMENTAL
/gene= "ORF"
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/product= "LlaAI GAIC
/evidence= EXPERIMENTAL
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Patent No. 6300109
Matches 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy
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COMPUTER: HAM BO comparible
OPERATING SYSTEM: PO DOS/MS DOS
SOFTWARE: Patentin malere
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                                                                                                                                                                                  OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: double
TOPOLORY: linear
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                                                                                                                                                                                                                                                NAME/KEY: CDS
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Pred. No. 16;
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Segmence 156, Application Pc/IUS91036H0
GENERAL INFORMATION:
APPLICANT: Mailtoneei, Mark D
APPLICANT: Krawczyk, Steven
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Hest Local Similarity [180.5%: Prod. No. 16]
Matches [9] Conservative [9] Mismatches of EdeCs
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                                                                   TITLE OF INVENTION: SEQUENCE-SPECIFIC WHICH EARLY CONTINUES OF INVENTION: SEQUENCE-SPECIFIC MARCHES HAS A LIBERTAL ACTION OF THE SEQUENCE SPECIFIC ACTION OF THE SECUENCE SPECIFICATION OF THE SECUENCE SPECIFICAT
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COMBUTER: IBM BY compatible

OPERATING SYSTEM. BY DVS/MS-108

SCETWARE: Patroith Release #1.0, Version #1.0 (EP);

OURRENT APPLICATION DATA:
CURRESPONDENCE ADDRESS
                                    NUMBER OF SEQUENCES:
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CTHER INFORMATION: /cob
OTHER INFORMATION: /cob
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MEDIUM LYPE: Floppy
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LENGTH: 369% base pairs
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TIBLE OF INVENTION: restriction modification systems from Lactococcus factio
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STRANDEDNESS: double
Tarolosky: linear
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ID-CALLON: 2412.
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/label- r IJaA:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM COMPATIBLE
COPERATING SYSTEM: PG-DGS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1 25
CURRENT APPLICATION DATA:
             ATTORNEY/AGENT INFORMATION
NAME: Seay, NICHOLAS J.
REGISTRATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: John, Maliyakal E.
TITLE OF INVENTION: TRANSGENIC COTTON PLANTS
TITLE OF INVENTION: PRODUCING HETEROLOGOUS PEROXIDASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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NAME: Murashige, Kate H.
REGISTRATION NUMBER: 29,
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                           STREET: P.O. B
CITY, Madison
STATE: WI
                                                                                                                                                                                                                                                                                                                              STREET: First Wisconsin Plaza, One South
STREET: Pinckney St.,
STREET: P.O. Box 2113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NÜMBER: 29,959
REFERENCE/DOCKET NUMBER: 4610-0011.40
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OPERATING SYSTEM:
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CITY: Menlo Park
                                                                                           CLASSIFICATION:
                                                                                                        FILING DATE:
                                                                                                                          APPLICATION NUMBER: US/08/378,588
                                                                                                                                                                                                                                                                       COUNTRY:
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SYSTEM: PC-DOS/MS-DOS
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TOPOLOGY: 11; MOLECULE TYPE: US-08-811-094-15
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                                                    Query Match
Best Local :
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Best Local :
                                   Matches
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INFORMATION FOR SEQ ID NO. SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                  TELEFAX: (608) 251-91 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                   FILING DATE: 25-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27,386
REFERENCE/DUCKET NUMBER: 11-2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECUT F ---
                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
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TITLE OF INVENTION: PRODUCING HETEROLOGOUS PEROXIDASE
NUMBER OF SEQUENCES: 29
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699 taattaattaattaatca 716
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                                   Local Similarity
hes 18, Conserv
                                                                                                                                                                           STRANDEDNESS: double
                                                                                                                                                                                            TYPE: nucleic acid
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P.O. Box 2113
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                                                                                                                                                                                                                                                                 : (608) 251-2484
(608) 251-9166
                               2.5%; Score 18; EB 2; Length 1064, ilarity 100.0%; Pred. No. 46; Conservative 0, Mishatches 0 Todels
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IBM PC compatible
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Matches 18; Conservative
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                                                                                                                                                                                                                             GENERAL INFORMATION:
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INFORMATION FOR SEQ 1D NO:
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REFERENCE/DOCKET NUMBER: 1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 251 2484
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LENGTH: 1064 base pair
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NAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Nichtlas ( Ser), Oberlos & Brit, STREET: First Wisconsin Plaza, one South
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                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Nicholas & Sear, Quarles & Brady
                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                       THEFT OF INVENTION:
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COMPUTER: HAM PC compatible
ODERATING SYSTEM - RC bass/MS bas
SCETWARE: Patentin Release #1.0, Version #1.25
                                 CITY: Madison
                                                     STREET: Pinckney St
STREET: P.O. Box 21
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COUNTRY
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                                                                                                                                                                     PRODUCTING HETEROLOGOUS PEROXIDASE
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Pred. No. 46;
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Best Local Similarity
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TELEFAX: (608) 251-9166
INFORMATION FOR SEQ ID NO: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: John, Maliyakal E.
TITLE OF INVENTION: TRANSGENIC COTTON PLANTS
TITLE OF INVENTION: PRODUCING HETEROLOGOUS PEROXIDASE
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                                    REFERENCE/DOCKET NUMBER: 11
                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/378,588
                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.29 CURRENT APPLICATION DATA:
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                                                                                                                FILING DATE: 25 JAN 1945
ATTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM: PC DOS/MS DOS
SOFTWARE: Patentin Release #1.0
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TYPE: nucleic acid
STRANDEDNESS: doubte
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COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-1XOS/MS-DUS
                                                                            REGISTRATION NUMBER:
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REGISTRATION NUMBER:
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                                                                                              Seay, Nicholas J
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P.O. Box 2113
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(608) 251 9166
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                    (608) 251 2484
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                                                                            27, 386
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APPLICATION:
CLASSIFICATION:
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
ANAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 11-229-9076-8
TELECOMMUNICATION INFORMATION:
TELEFAX: (608) 251-2484
TELEPHONE: (608) 251-2484
TELEFAX: (6
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Search completed: September 1, 2002, 03:12:46
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PCT-US94-11121-23
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APPLICANT: John, Maliyakal E.
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                                                                                                                                                                                                                                                                                                                                  Cuery Match 2.5%; Score 18; DB 5; Length 1270; Rest Local Similarity 100.0%; Pred. No. 45; Matches 18; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SUCTWARE PALENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/11121
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORPESSORE: Nicholas J. Seay, Quarles & Brady
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TITLE OF INVENTION: PRODUCING HETEROLOGOUS PEROXIDASE
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991 TAATTAATTAATTAATCA 1008
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TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
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Job time: 8235 sec